

FIG._1A

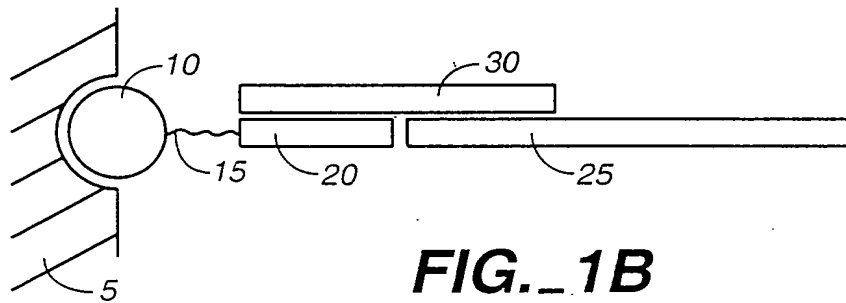


FIG._1B

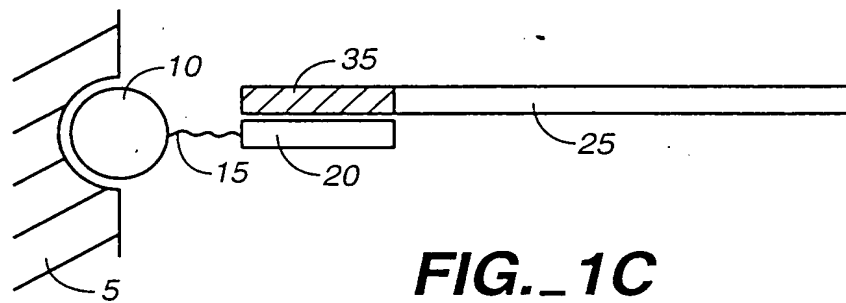


FIG._1C

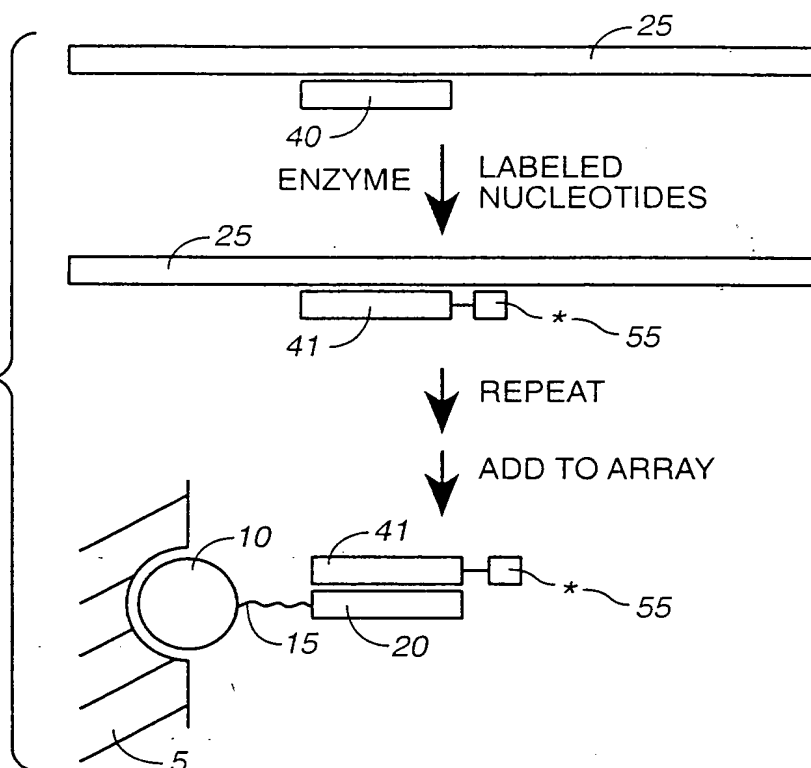
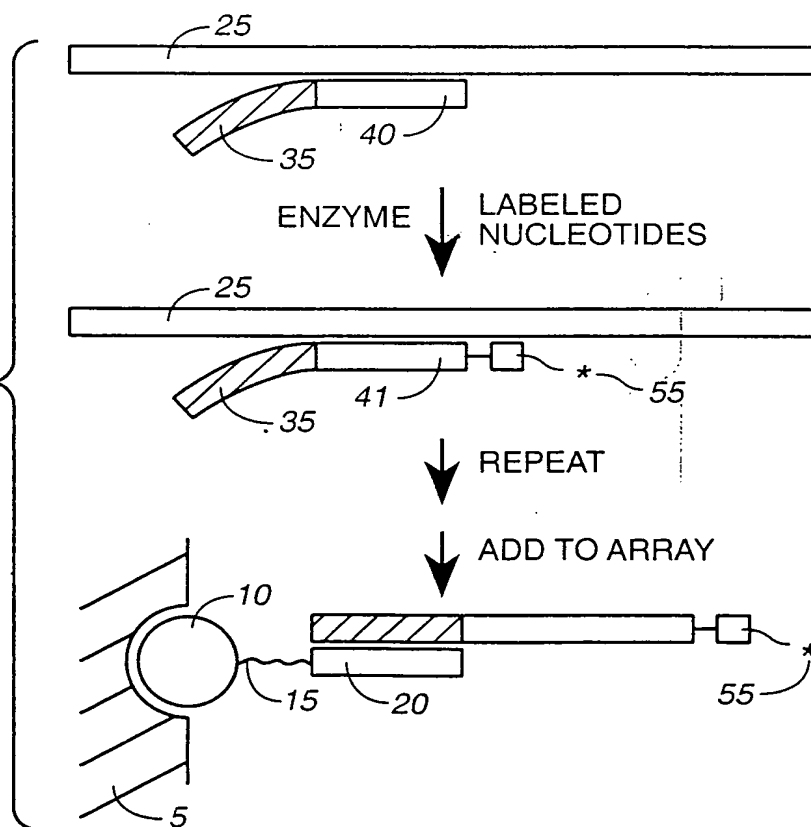
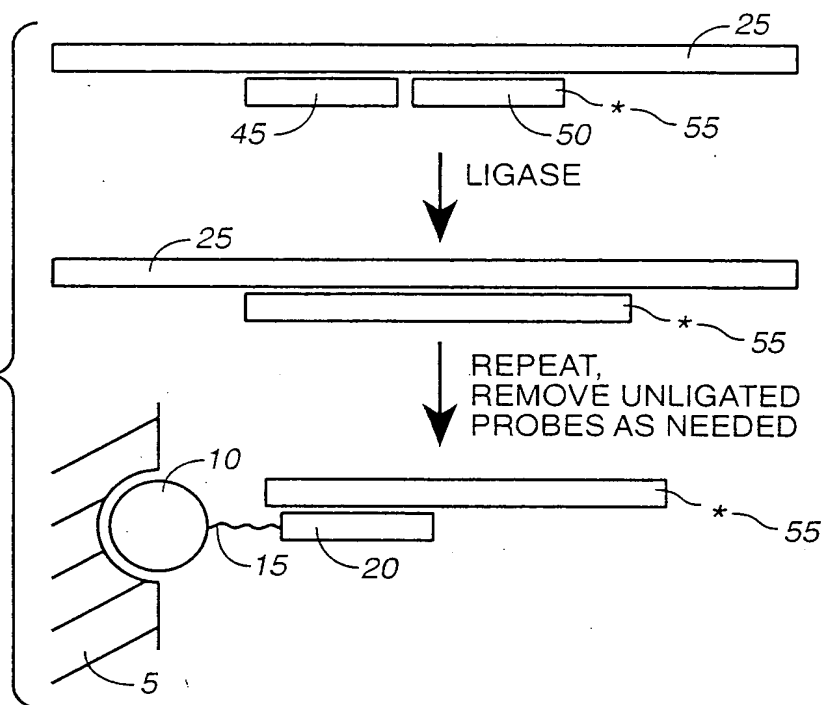
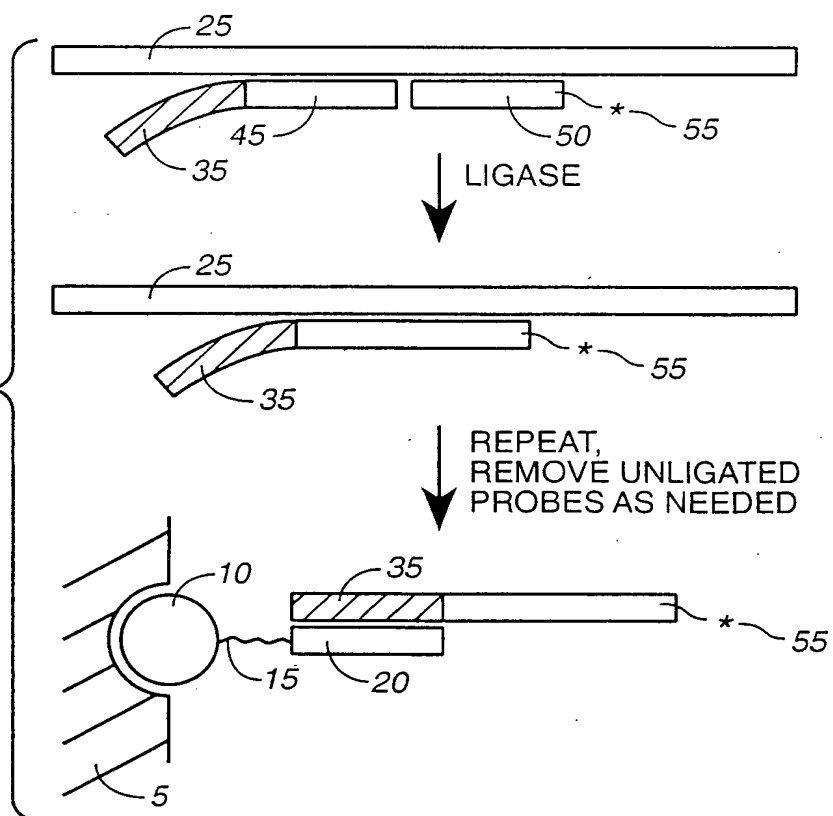
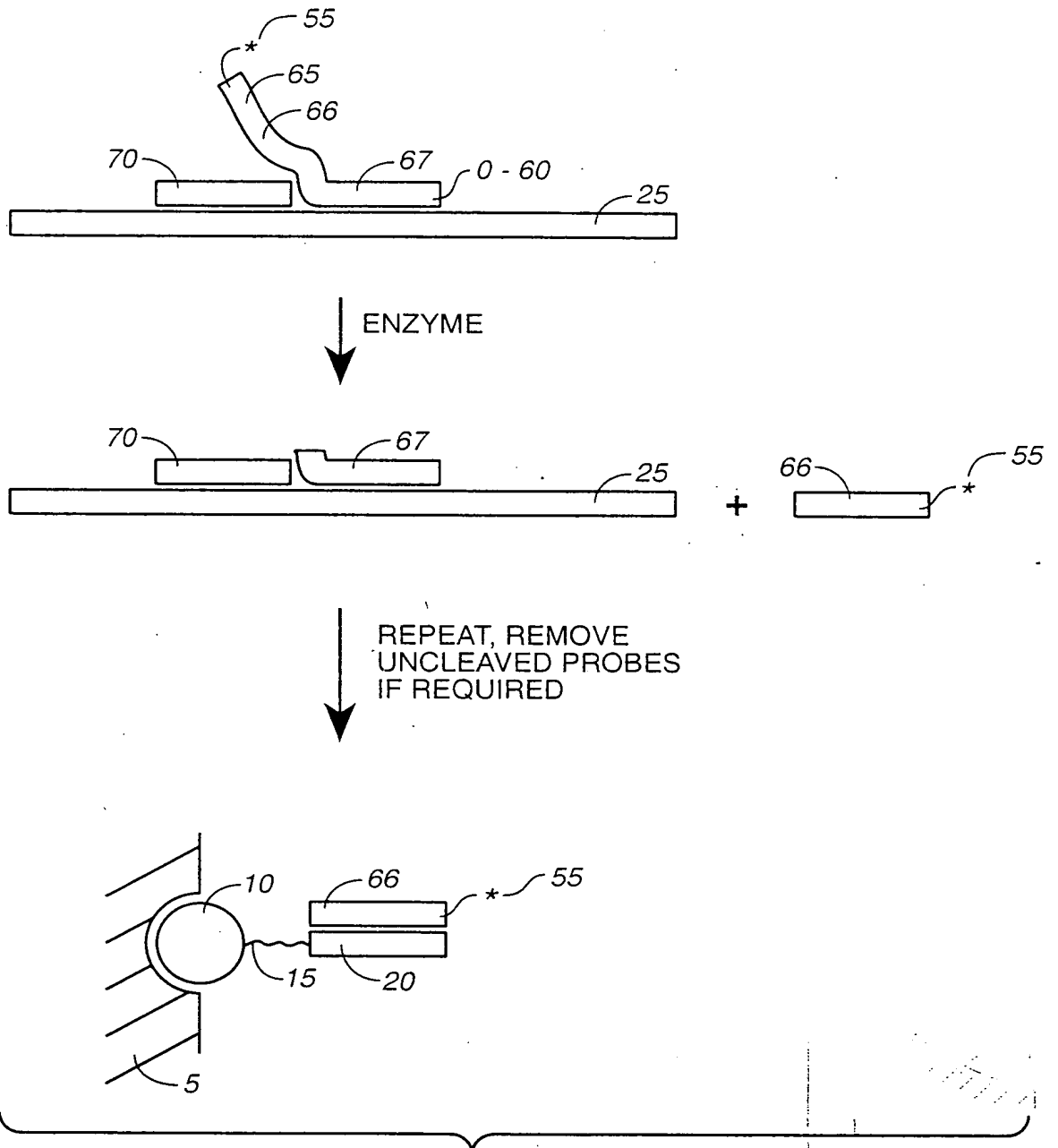
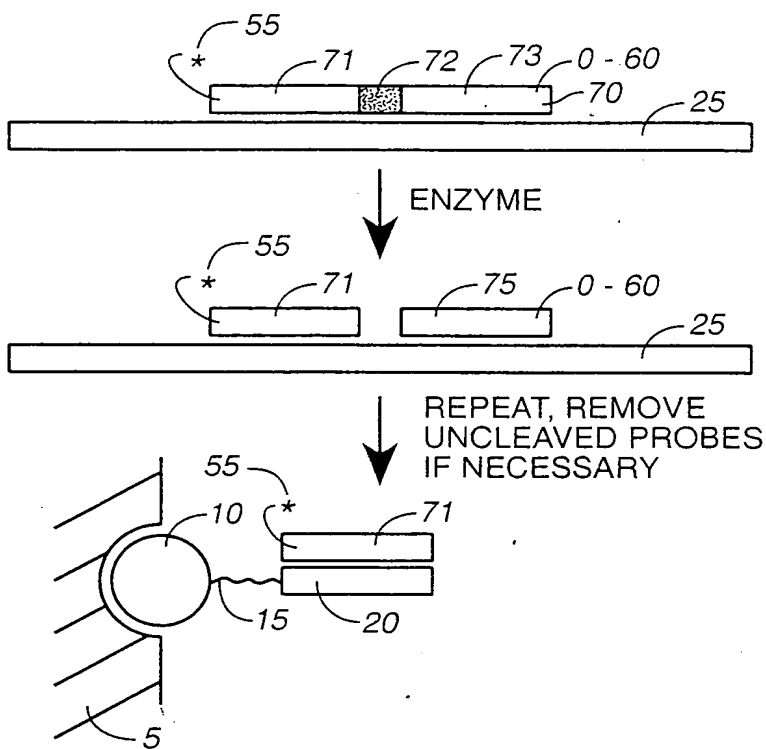
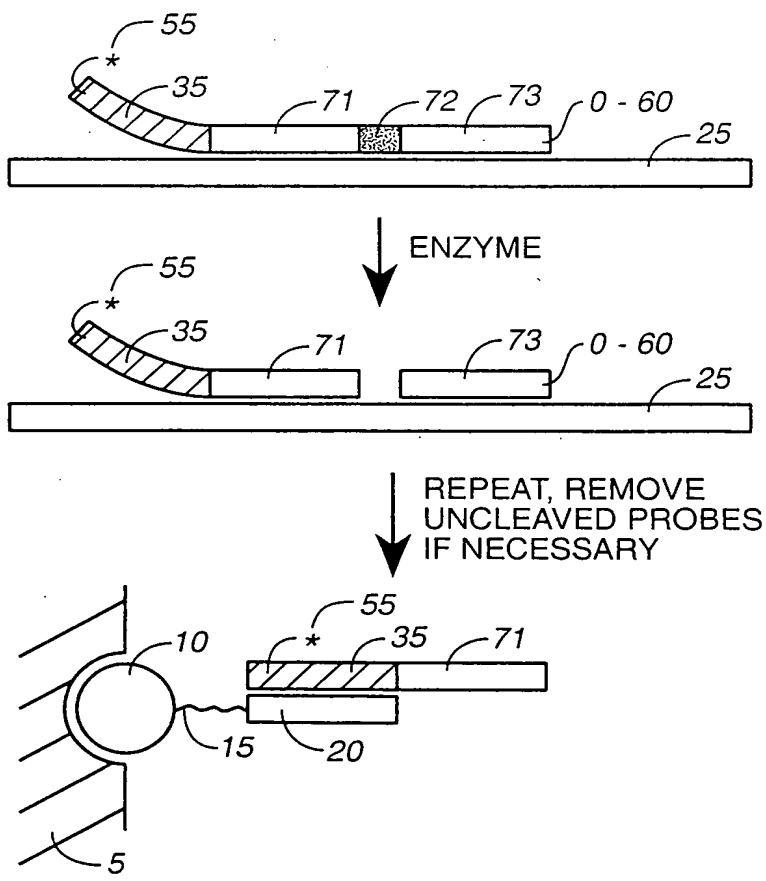
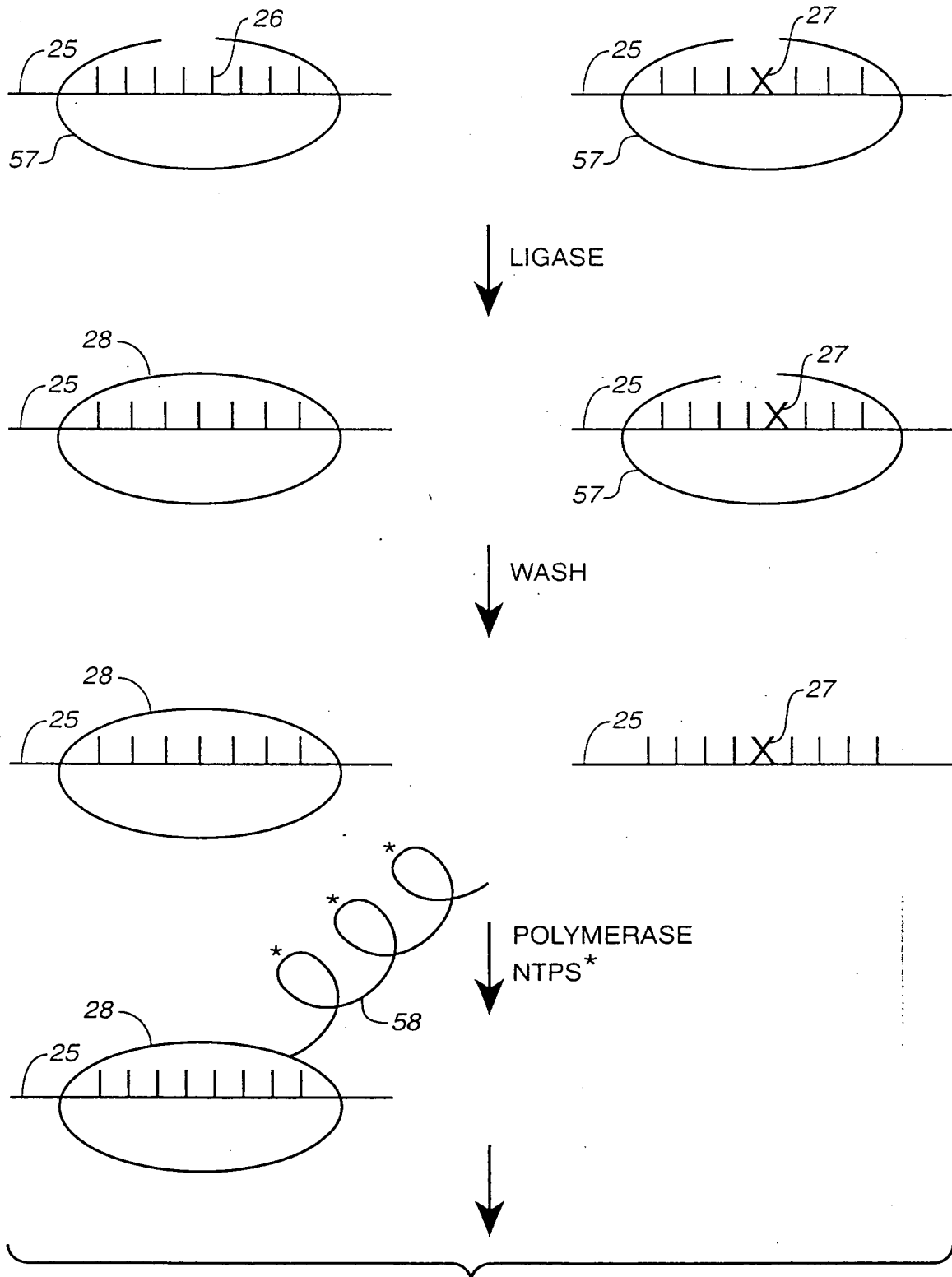
FIG._2A**FIG._2B**

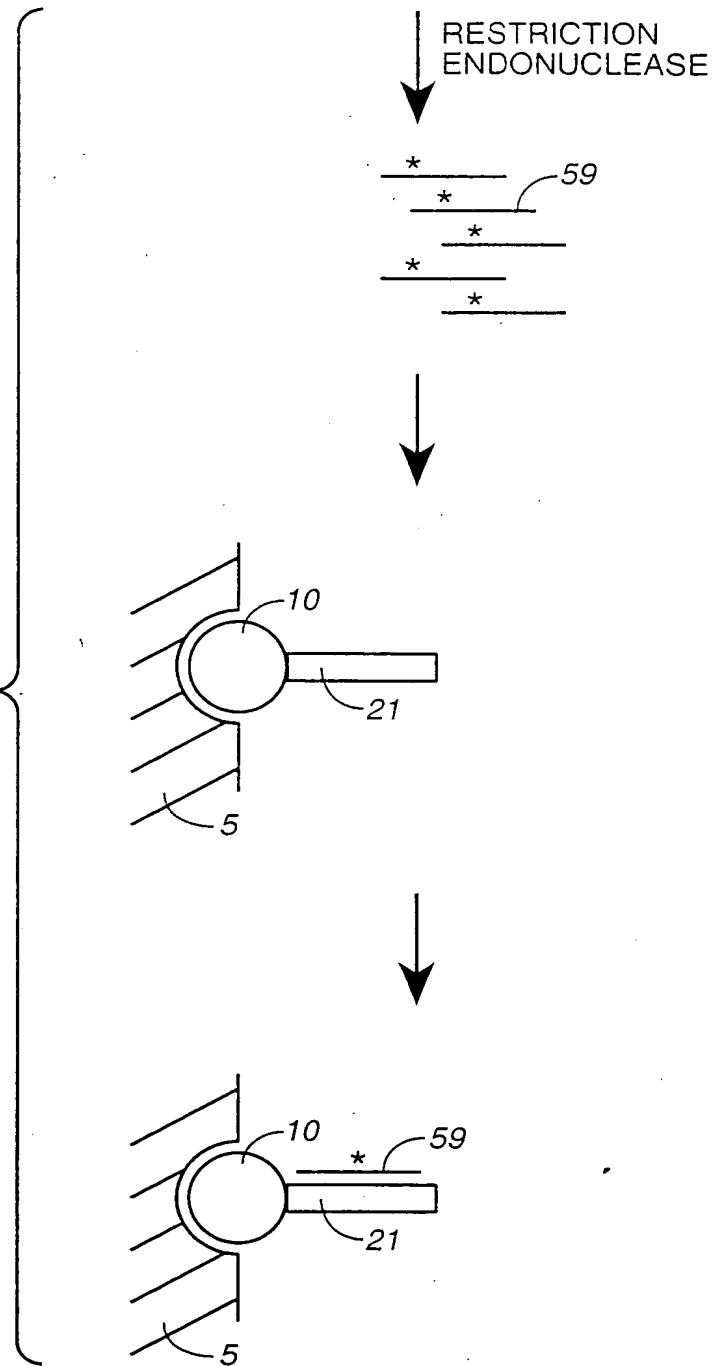
FIG._3A**FIG._3B**

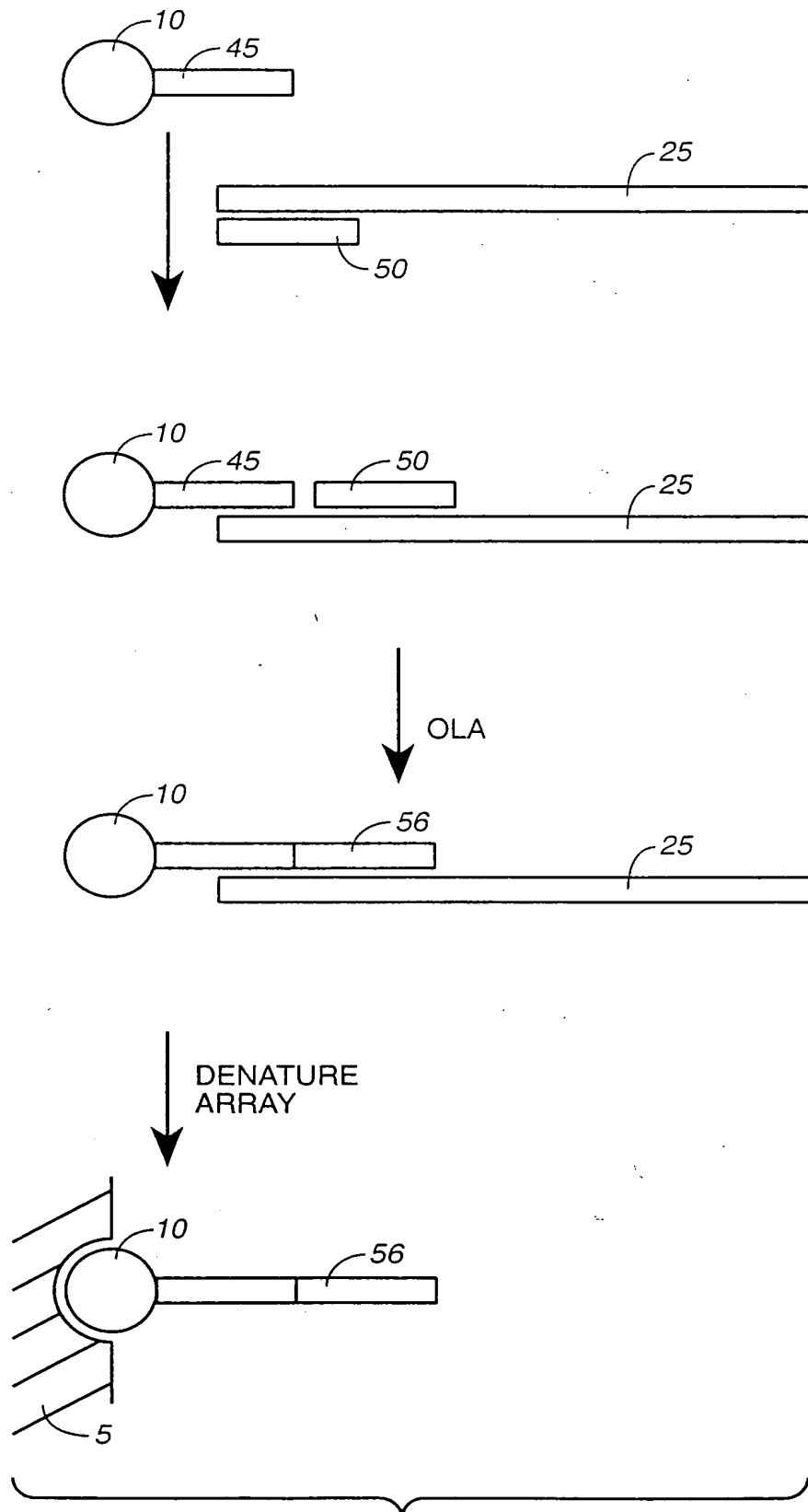
**FIG. 4**

5 / 36

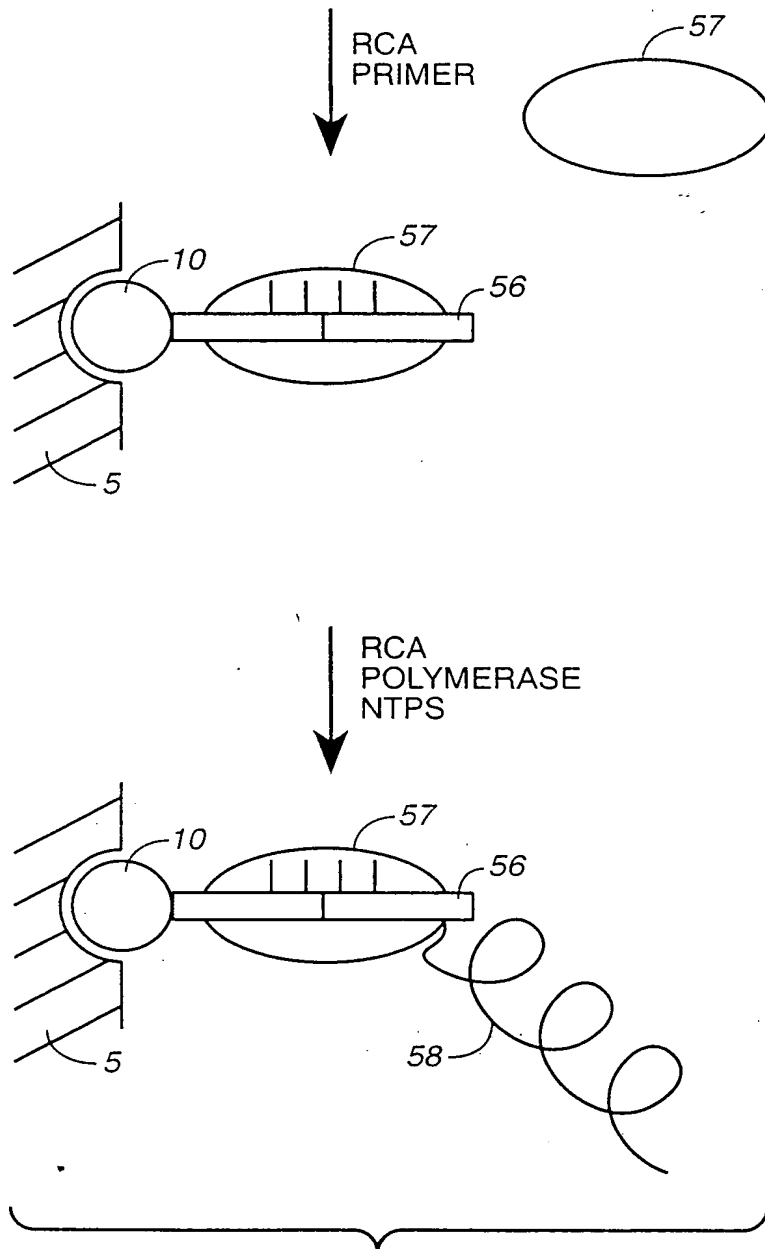
FIG. 5A**FIG. 5B**

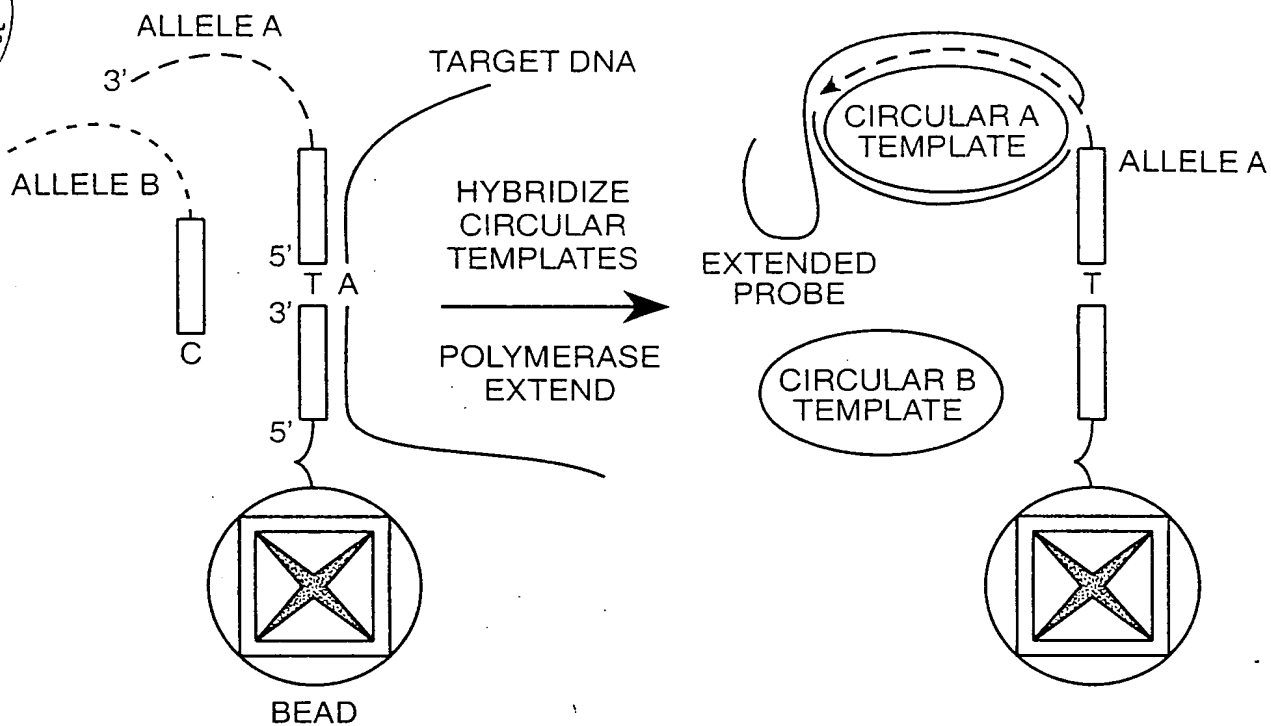
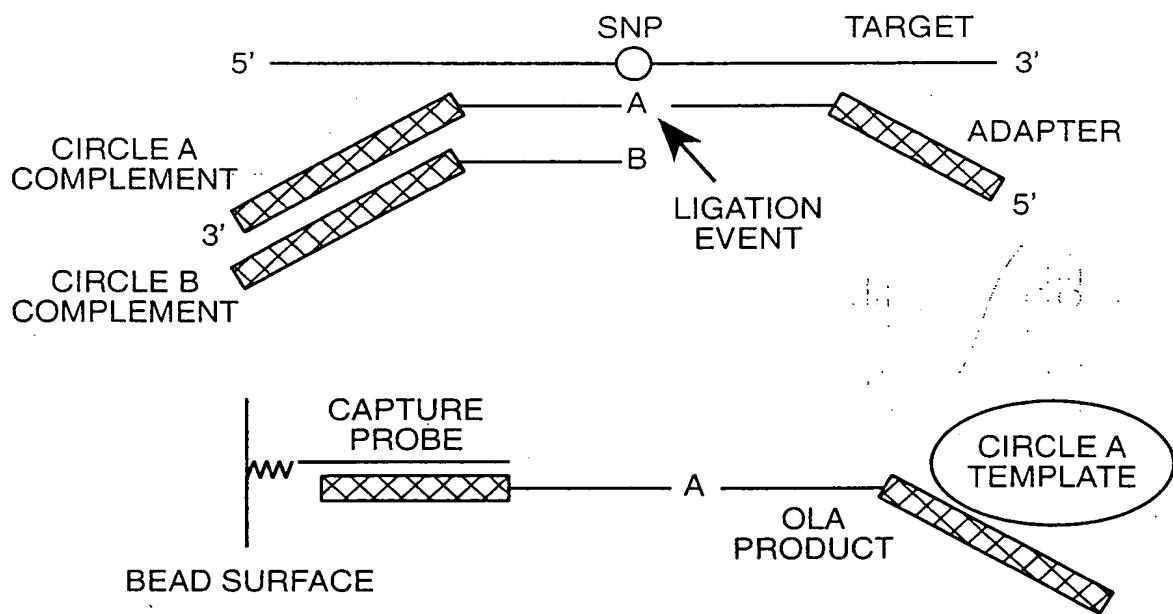
**FIG. 6A**

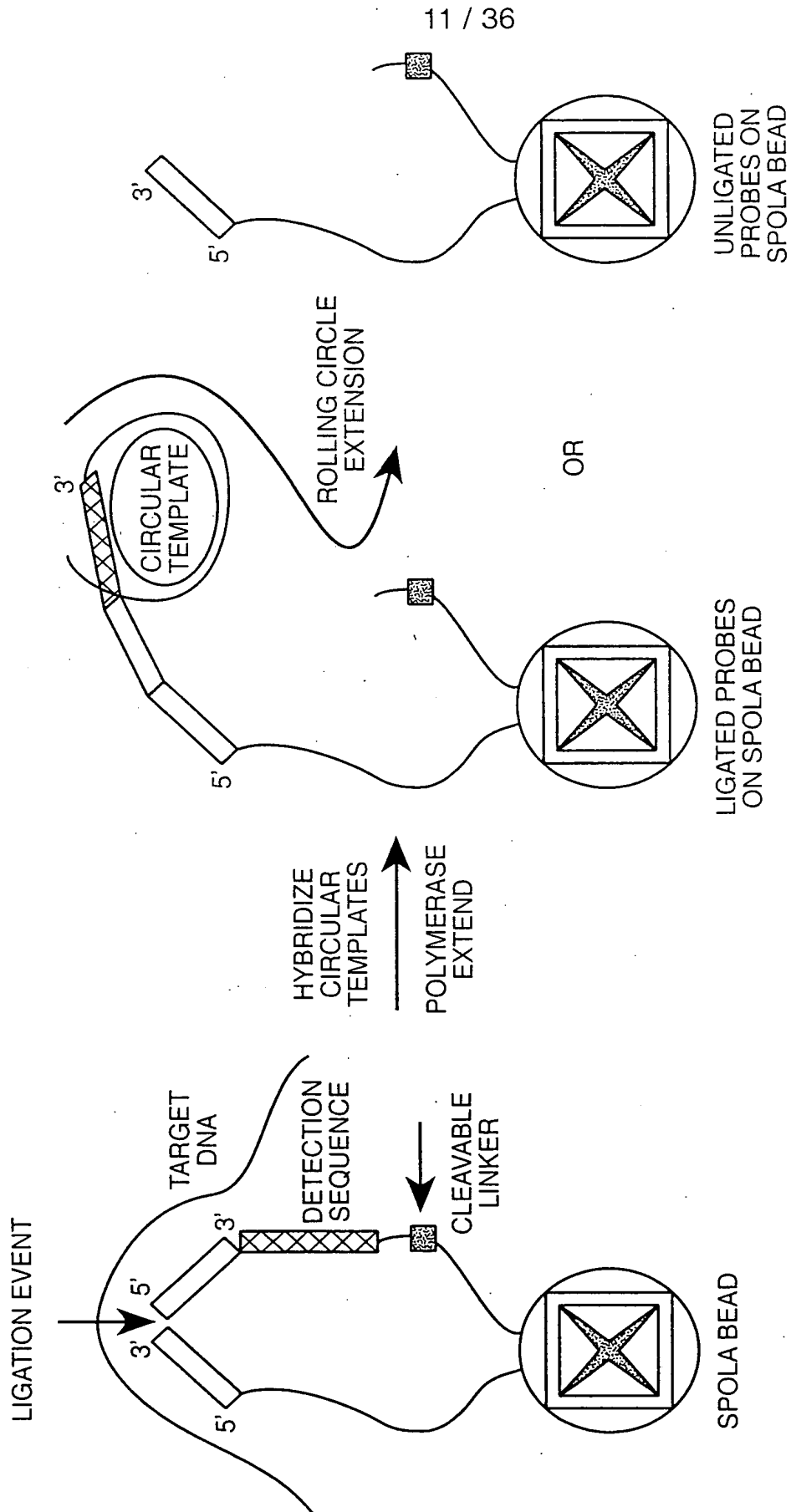
**FIG. 6B**

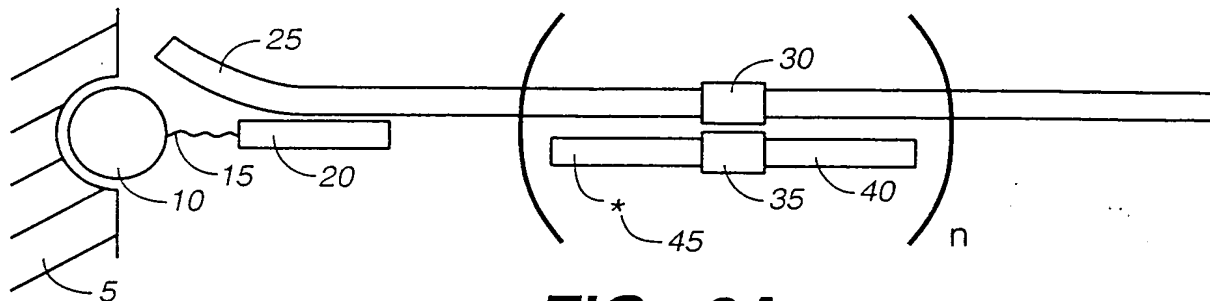
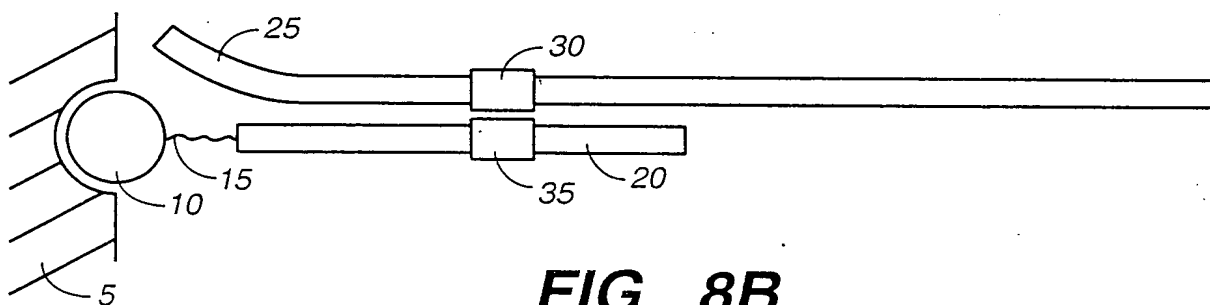
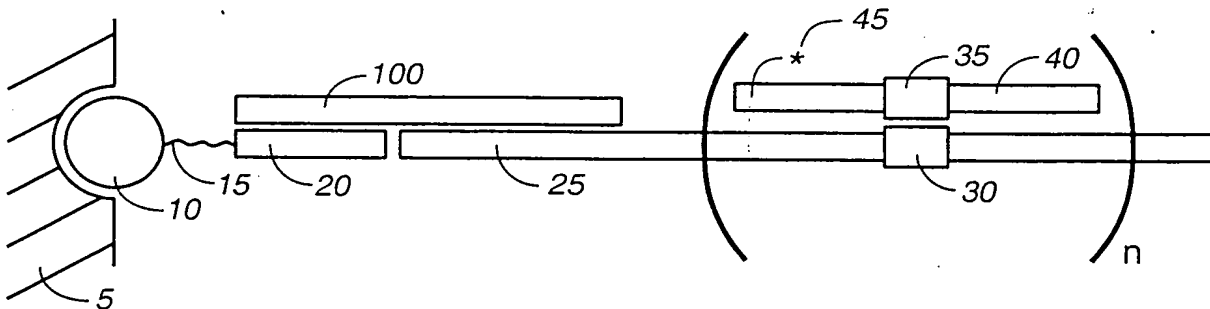
**FIG._7A-1**

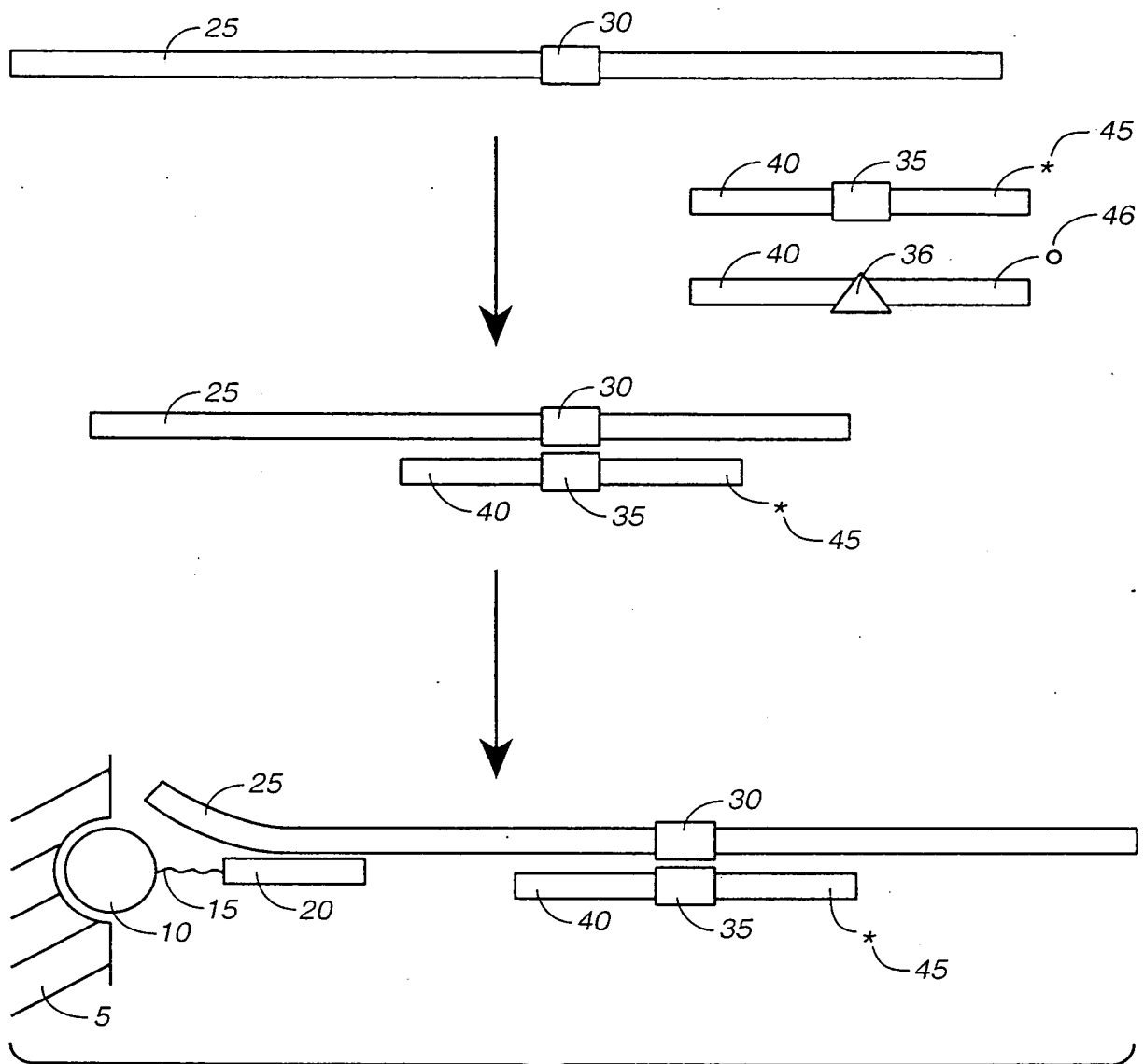
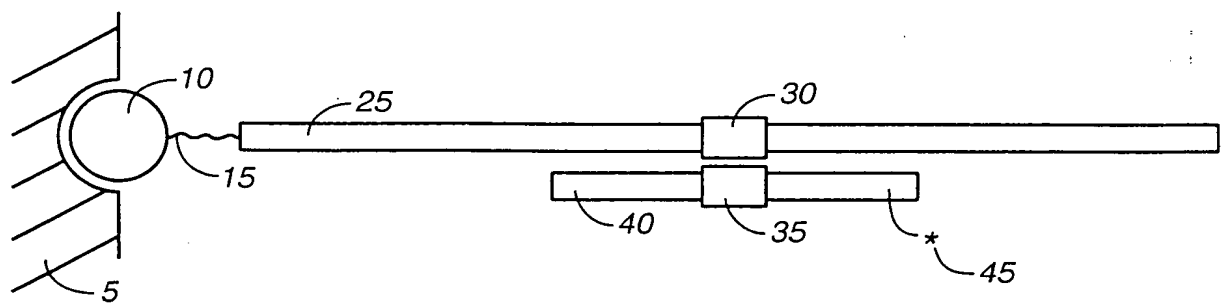
9 / 36

**FIG._7A-2**

**FIG._7B****FIG._7D**

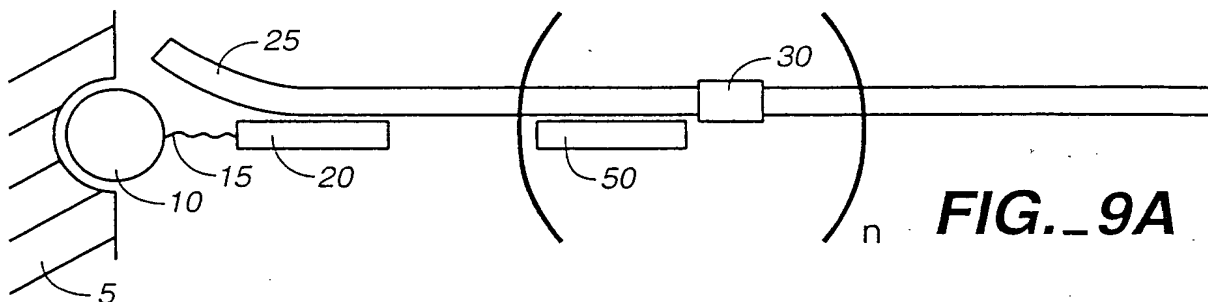
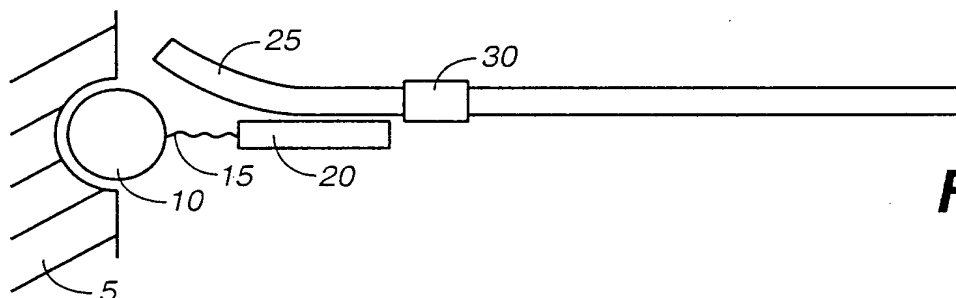
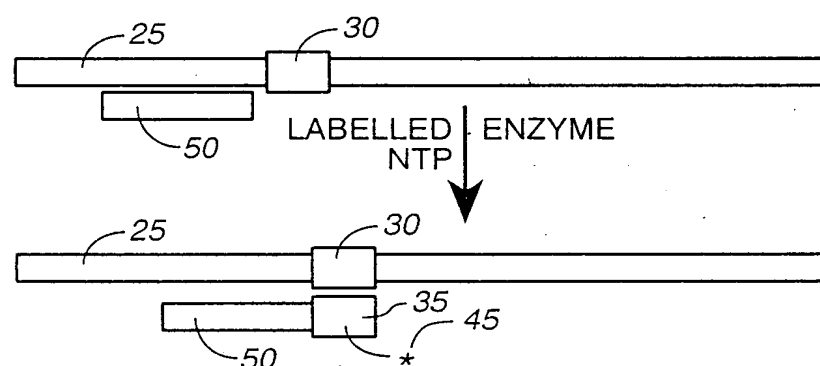


**FIG. 8A****FIG. 8B****FIG. 8C**

**FIG. 8D****FIG. 8E**

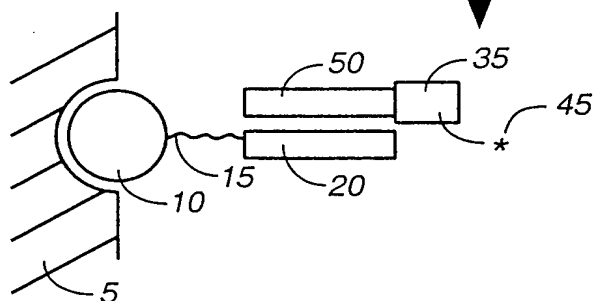
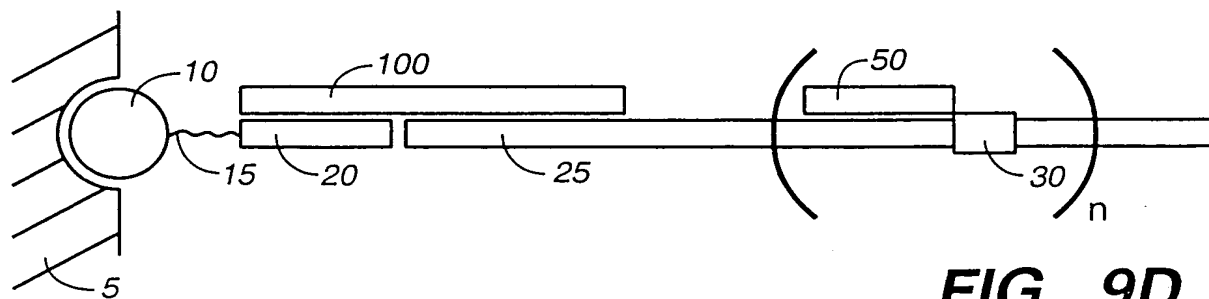


14 / 36

**FIG. 9A****FIG. 9B**

OPTIONAL REMOVAL OF
UNEXTENDED PRIMERS

DENATURE,
ADD TO ARRAY

**FIG. 9C****FIG. 9D**

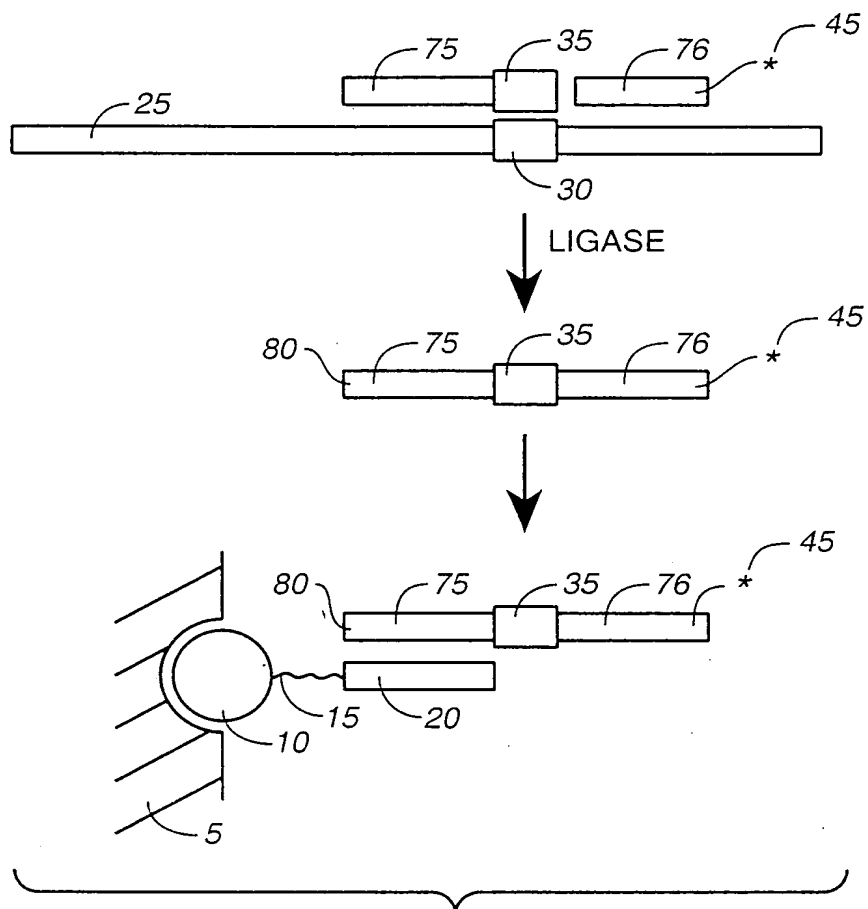


FIG._10A

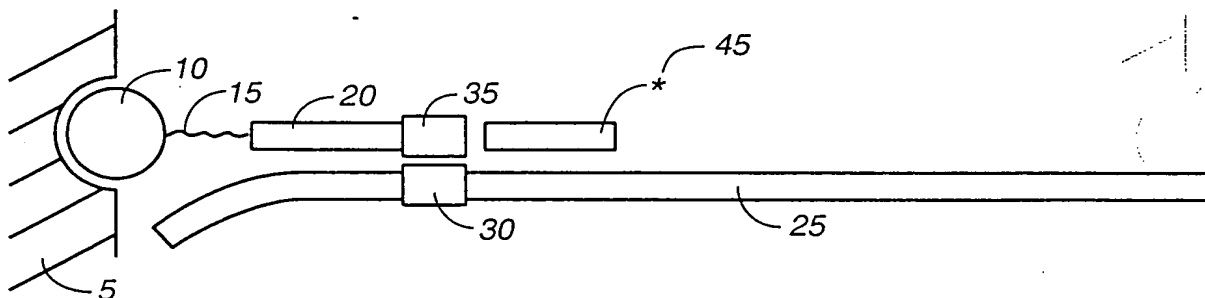
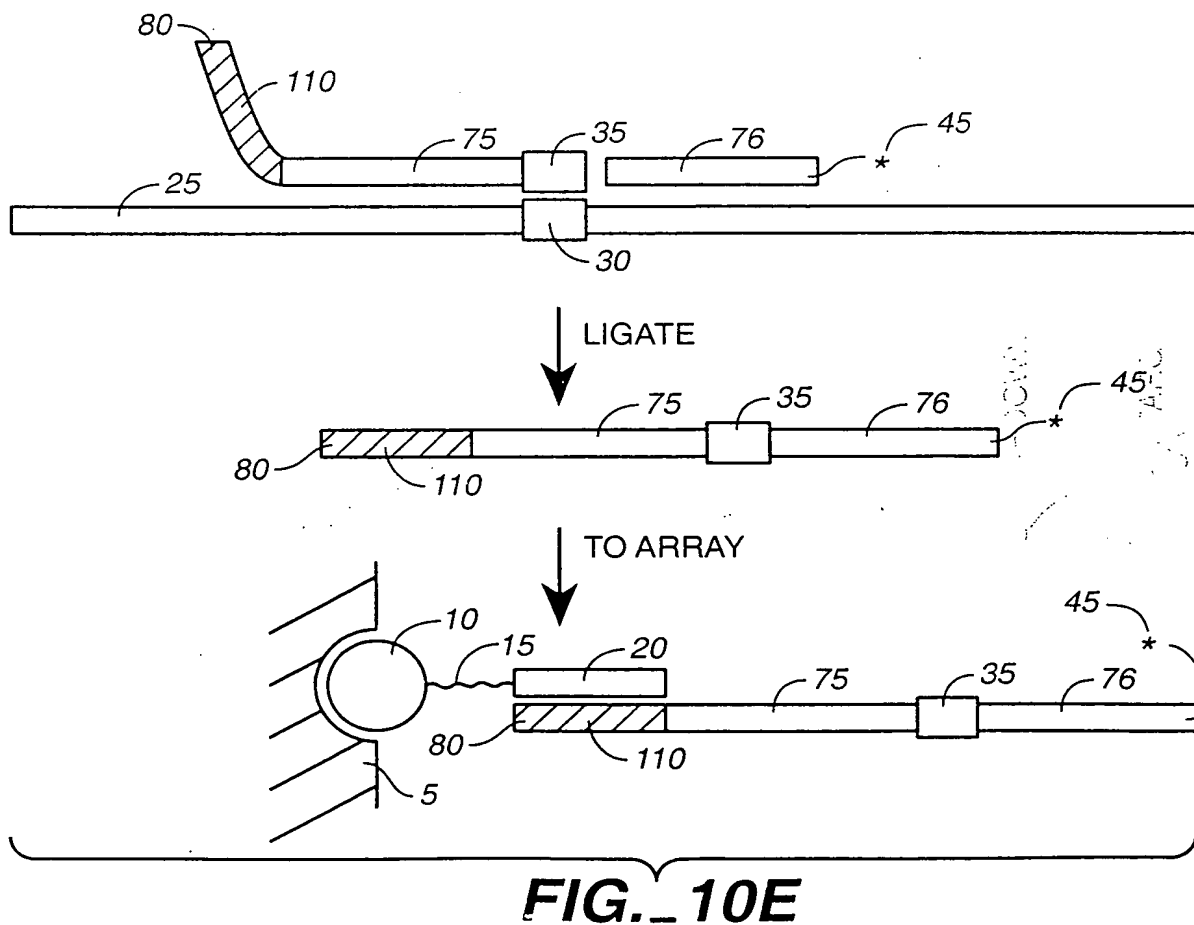
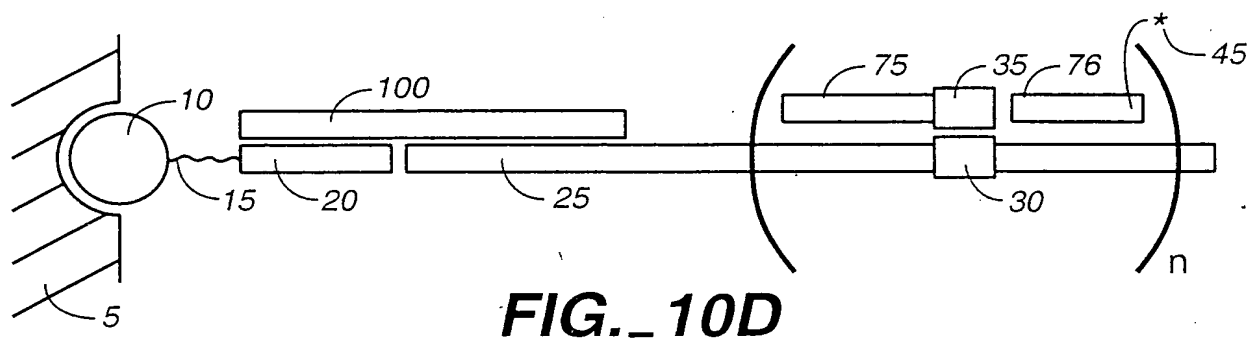
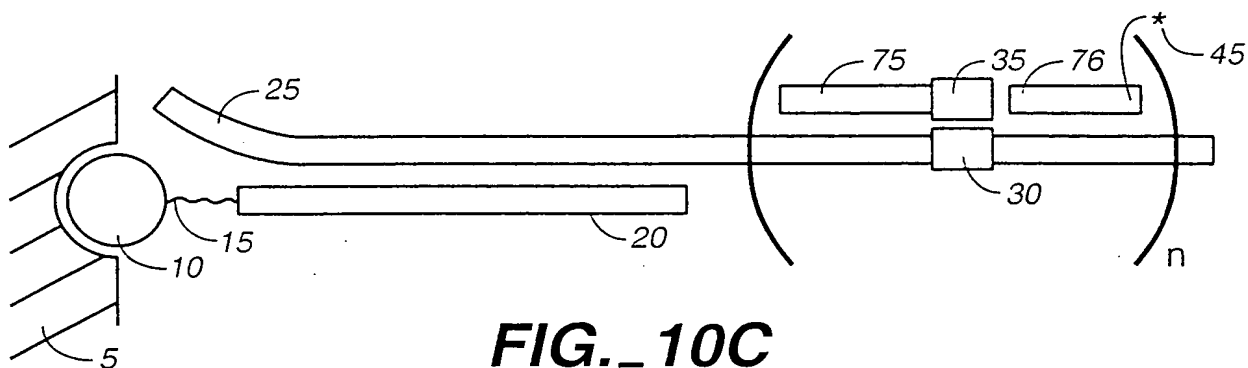


FIG._10B



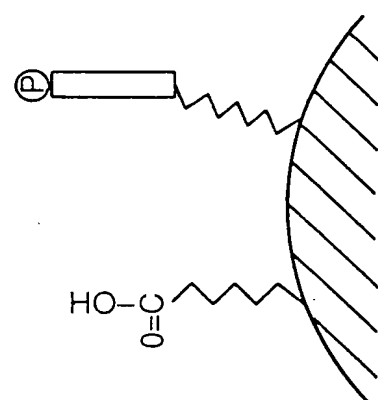
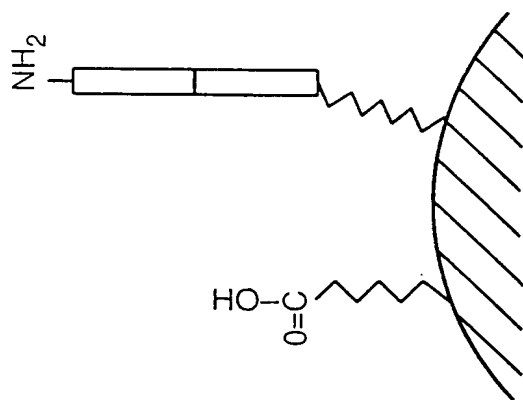


FIG. 11C

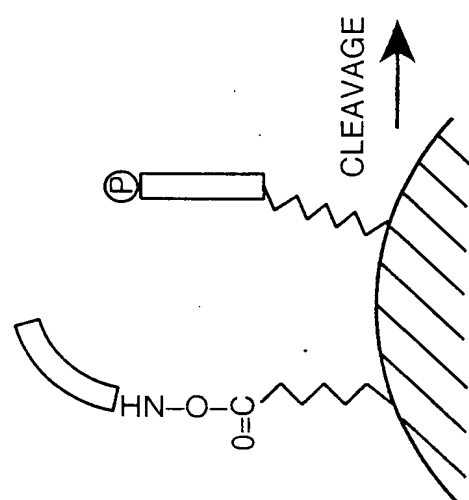
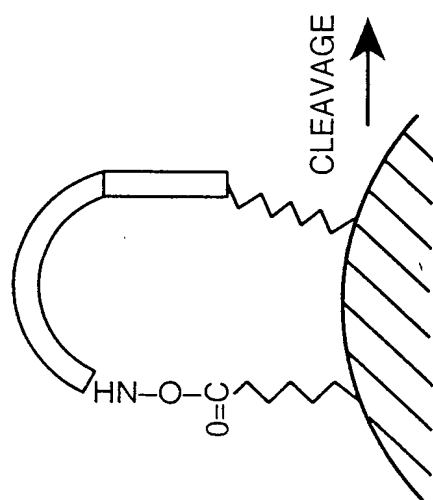


FIG. 11B

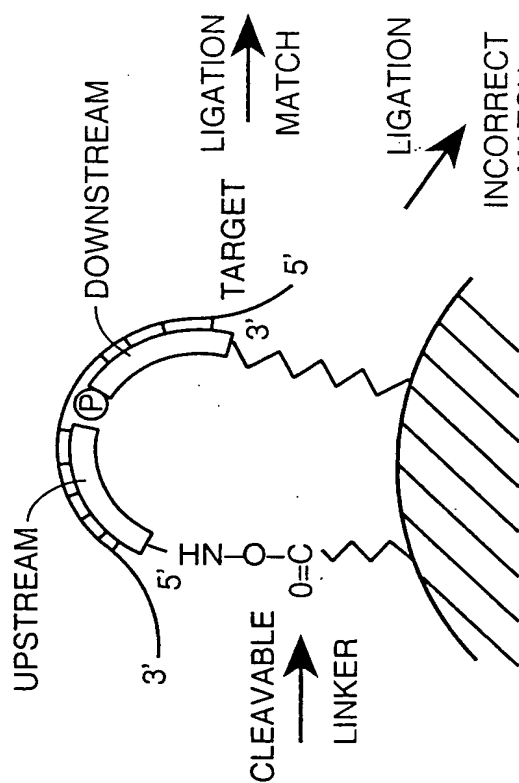
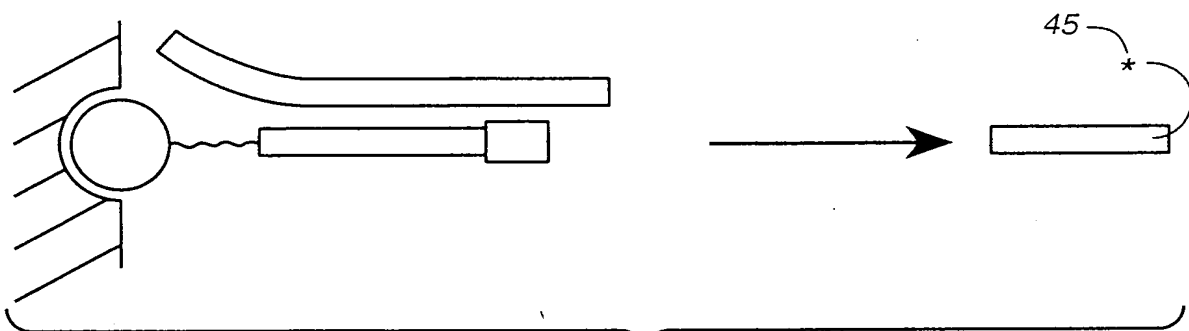
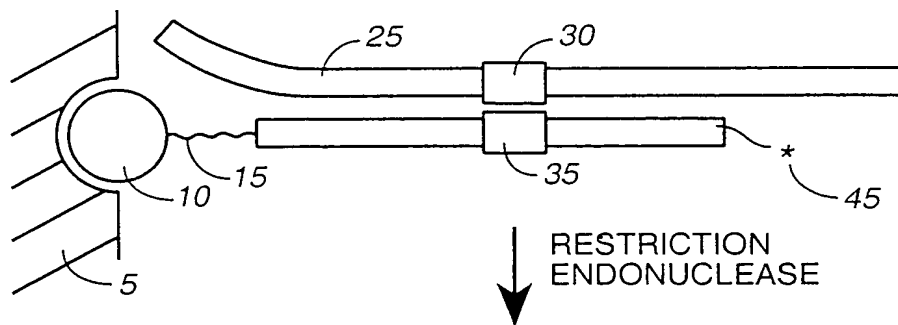
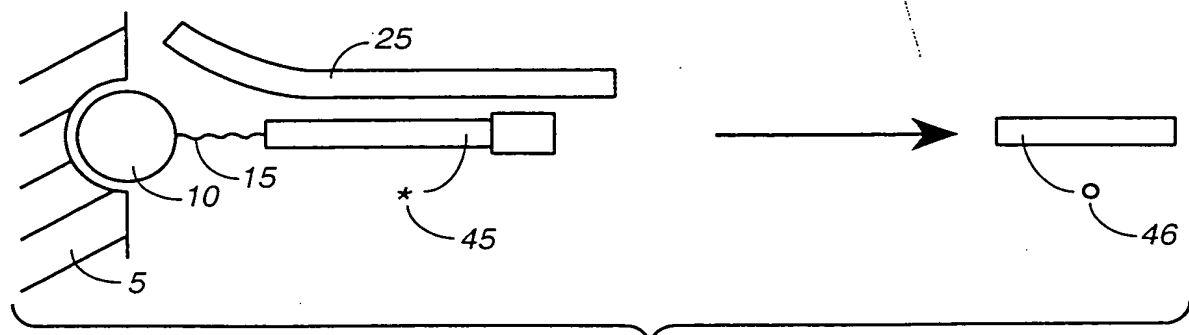
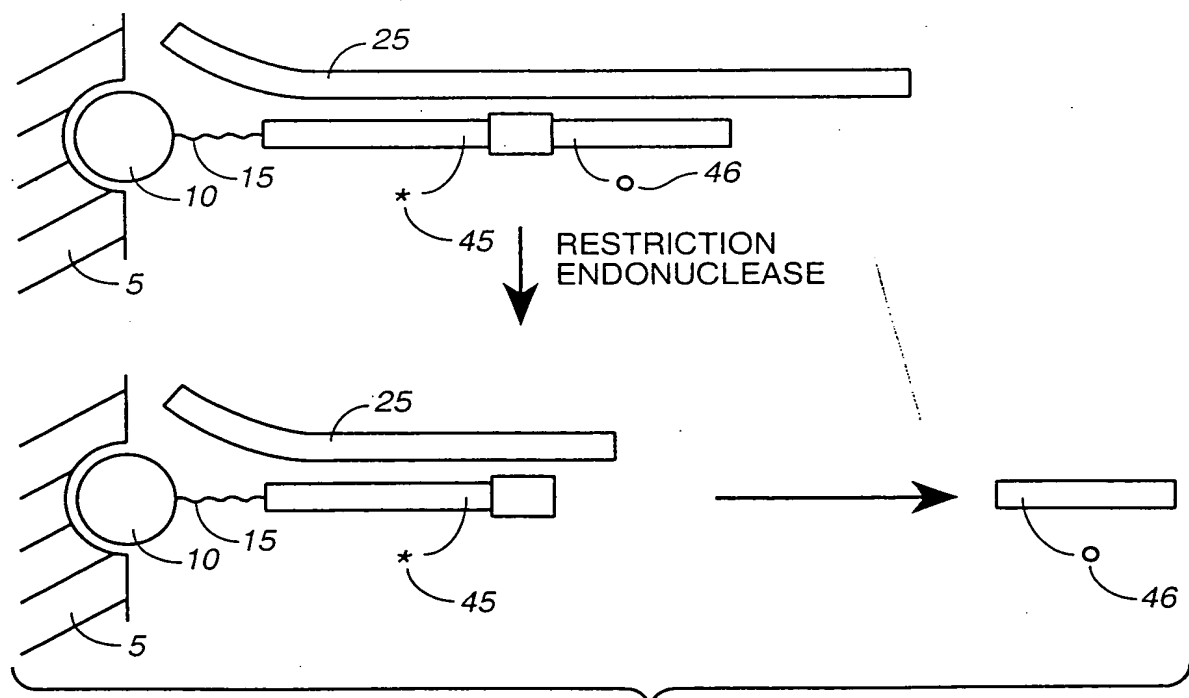
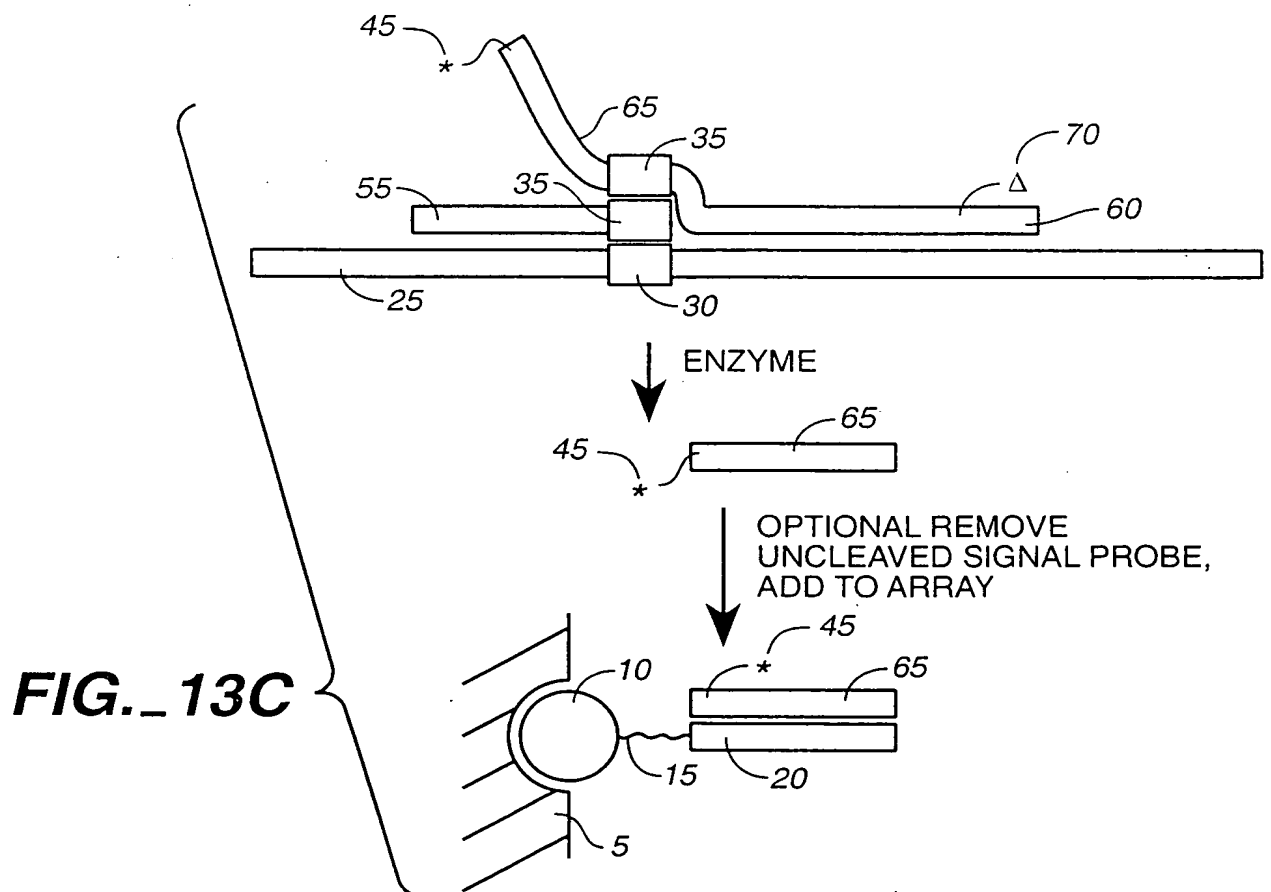
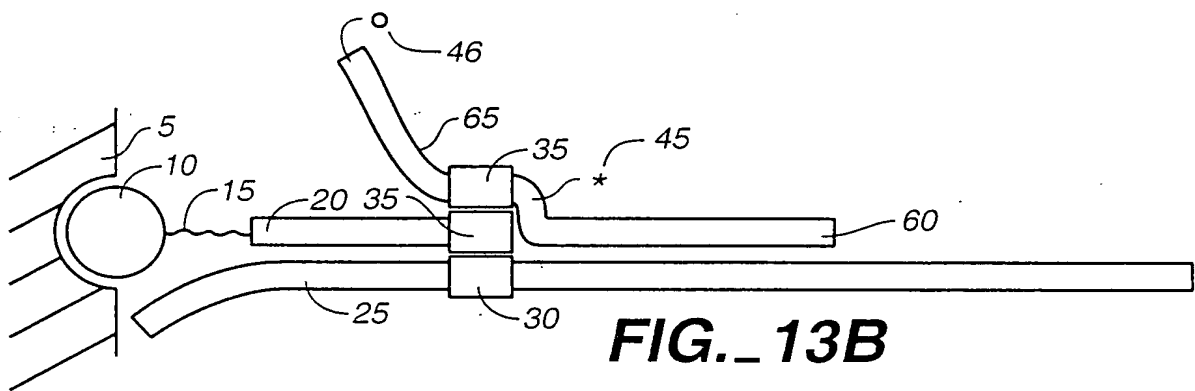
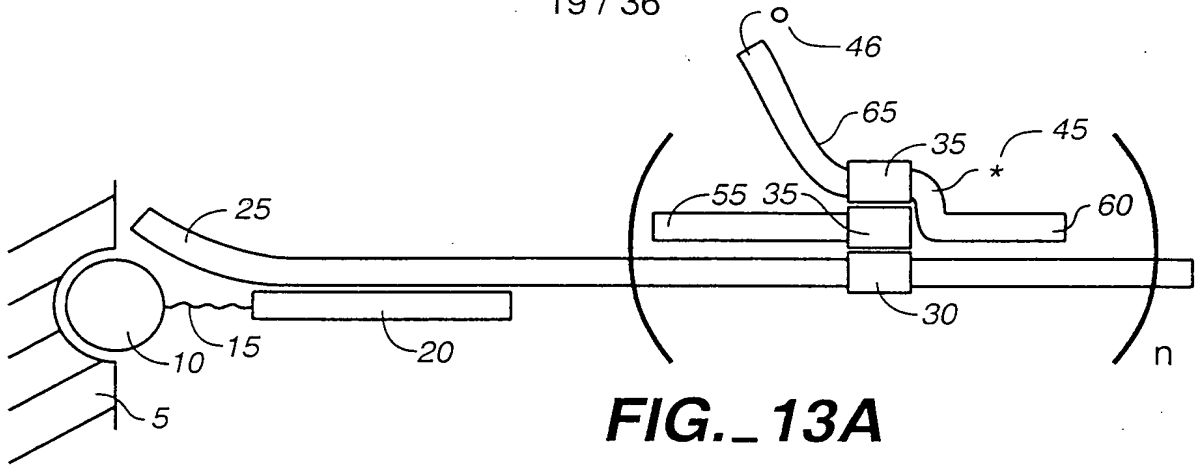


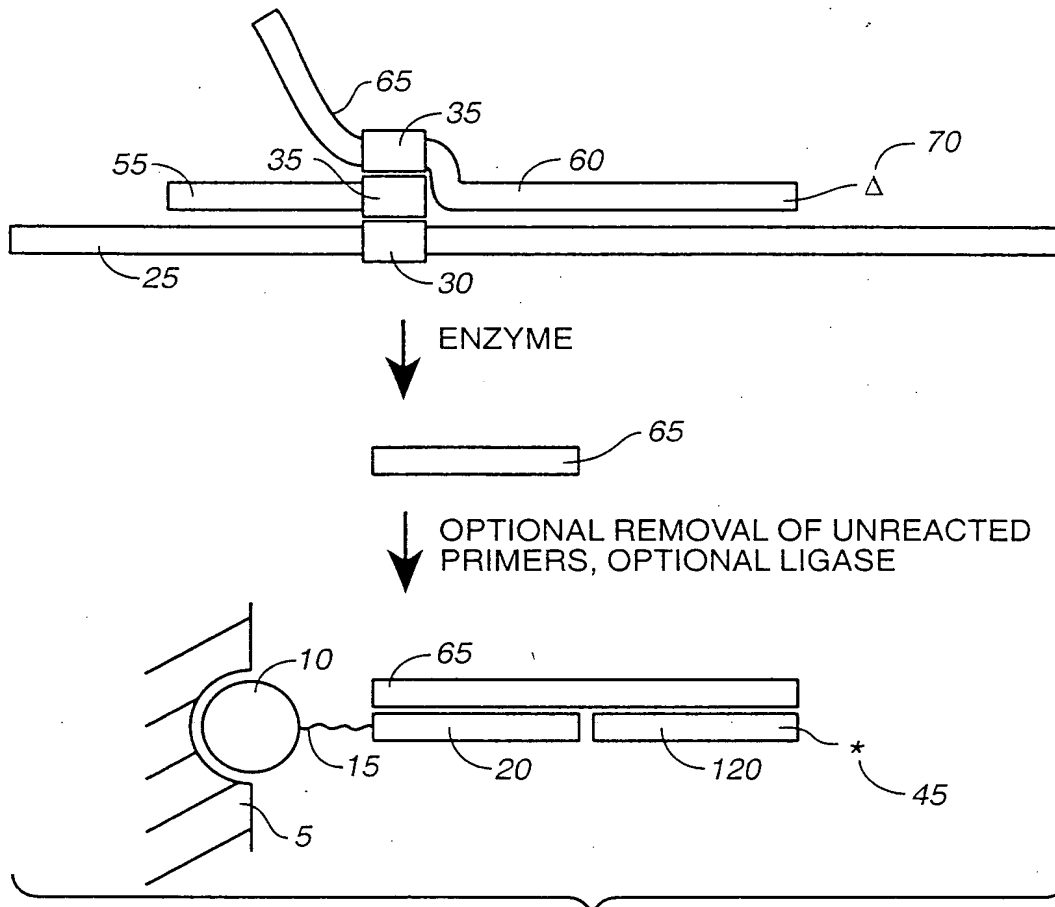
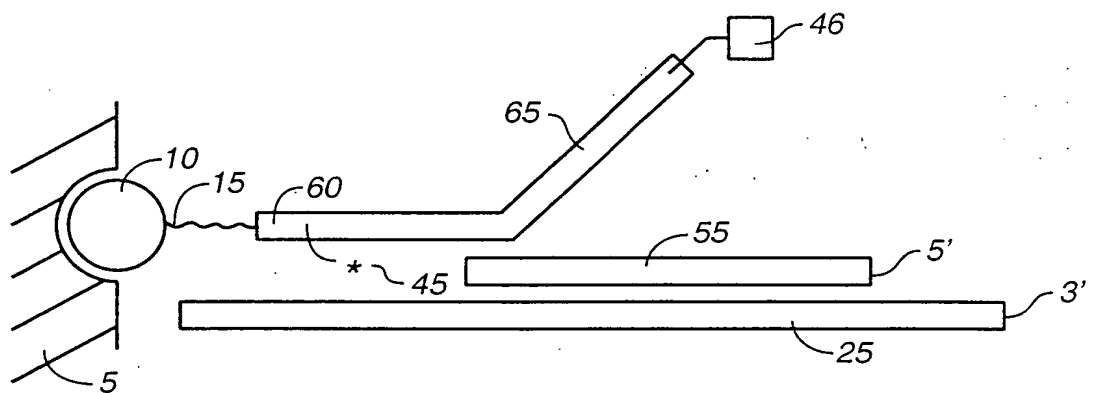
FIG. 11A

**FIG. 12A****FIG. 12B**



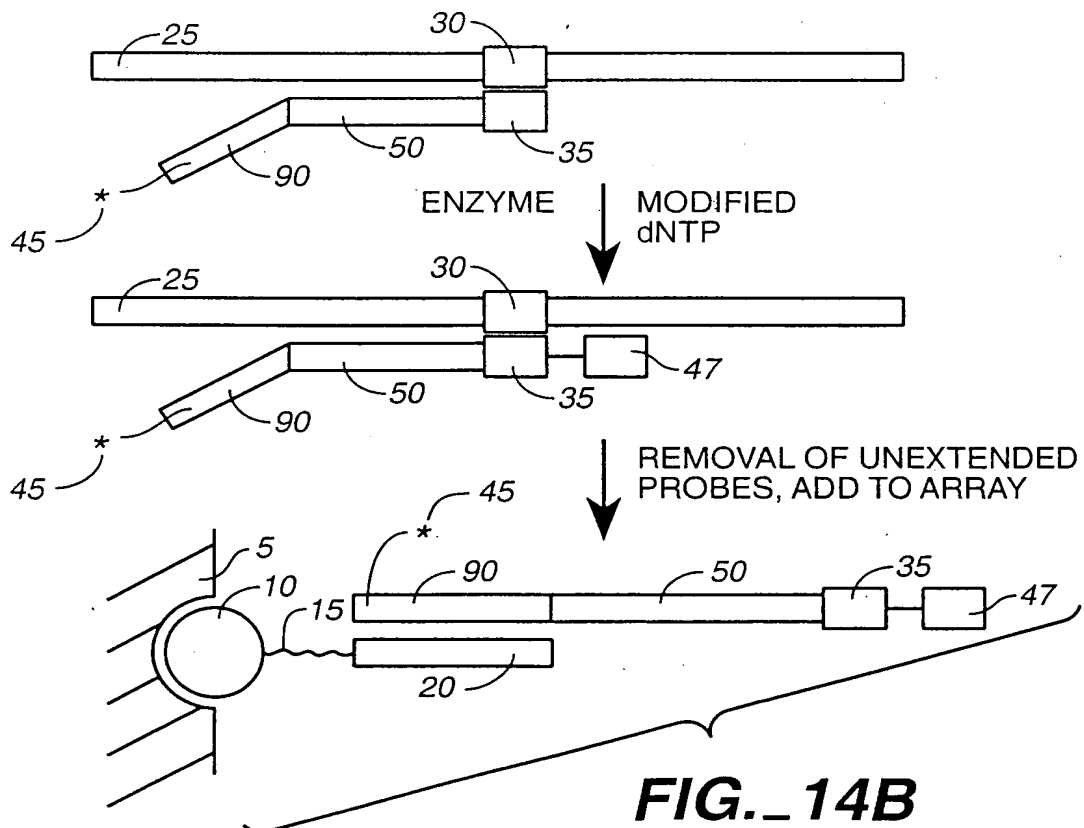
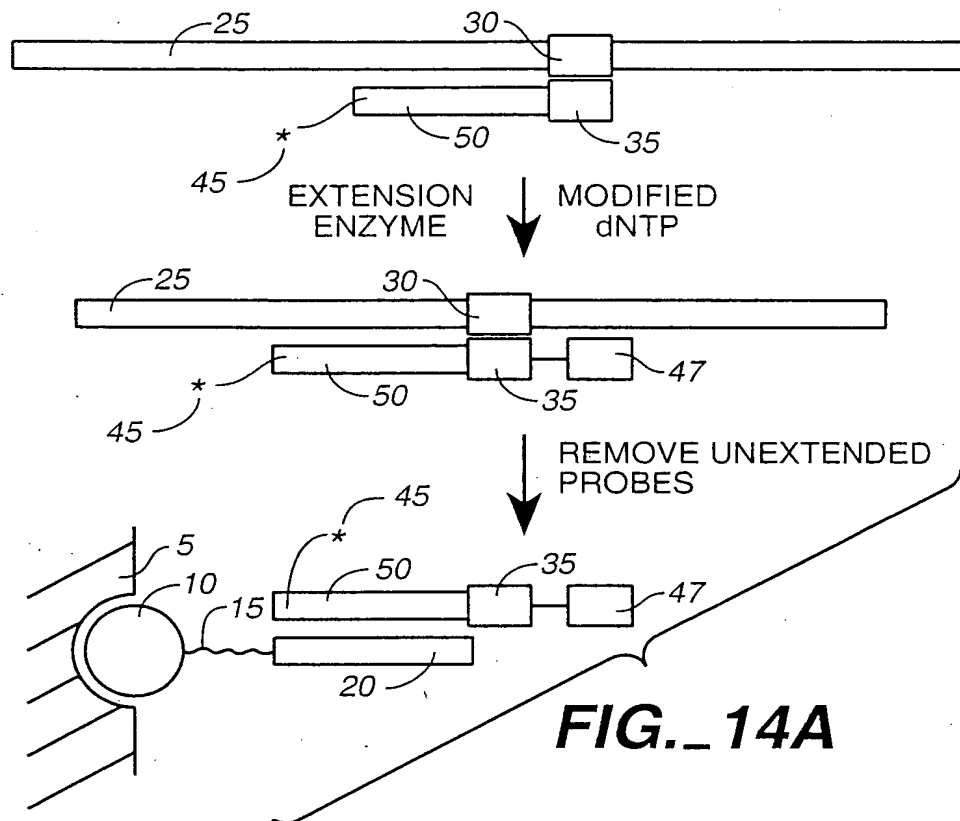
19 / 36

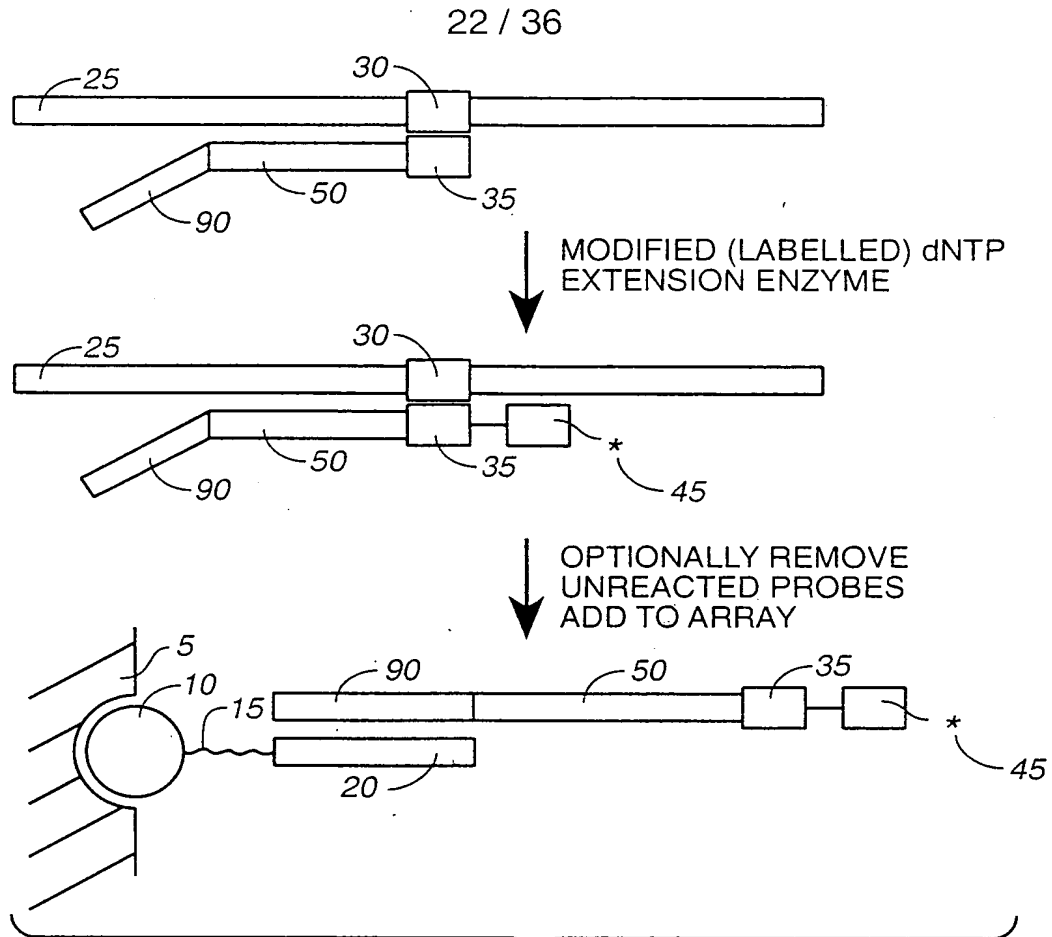
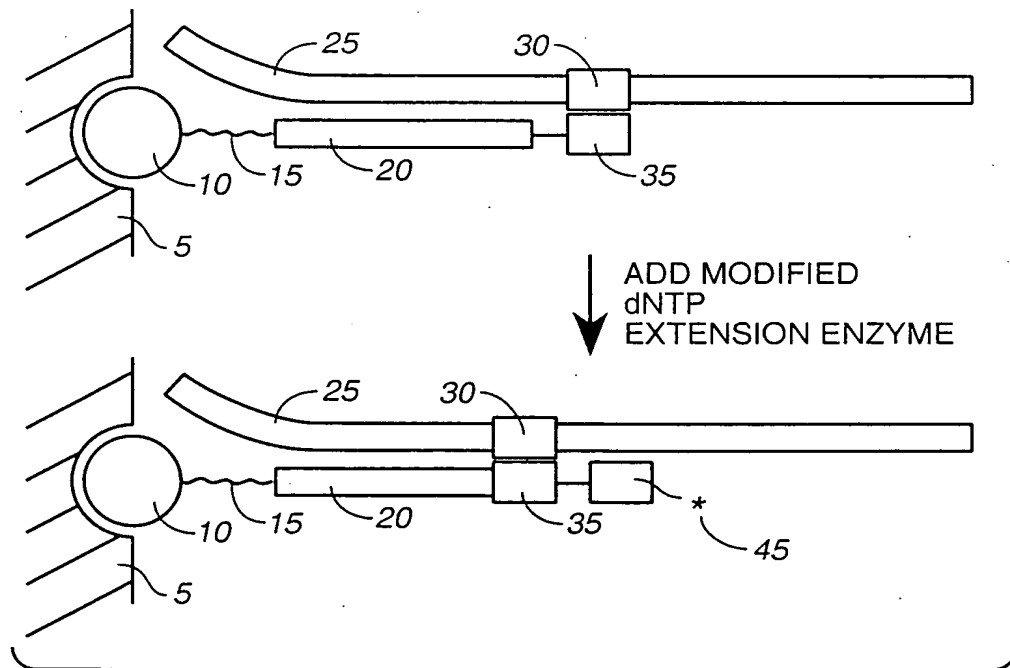


**FIG. 13D****FIG. 13E**

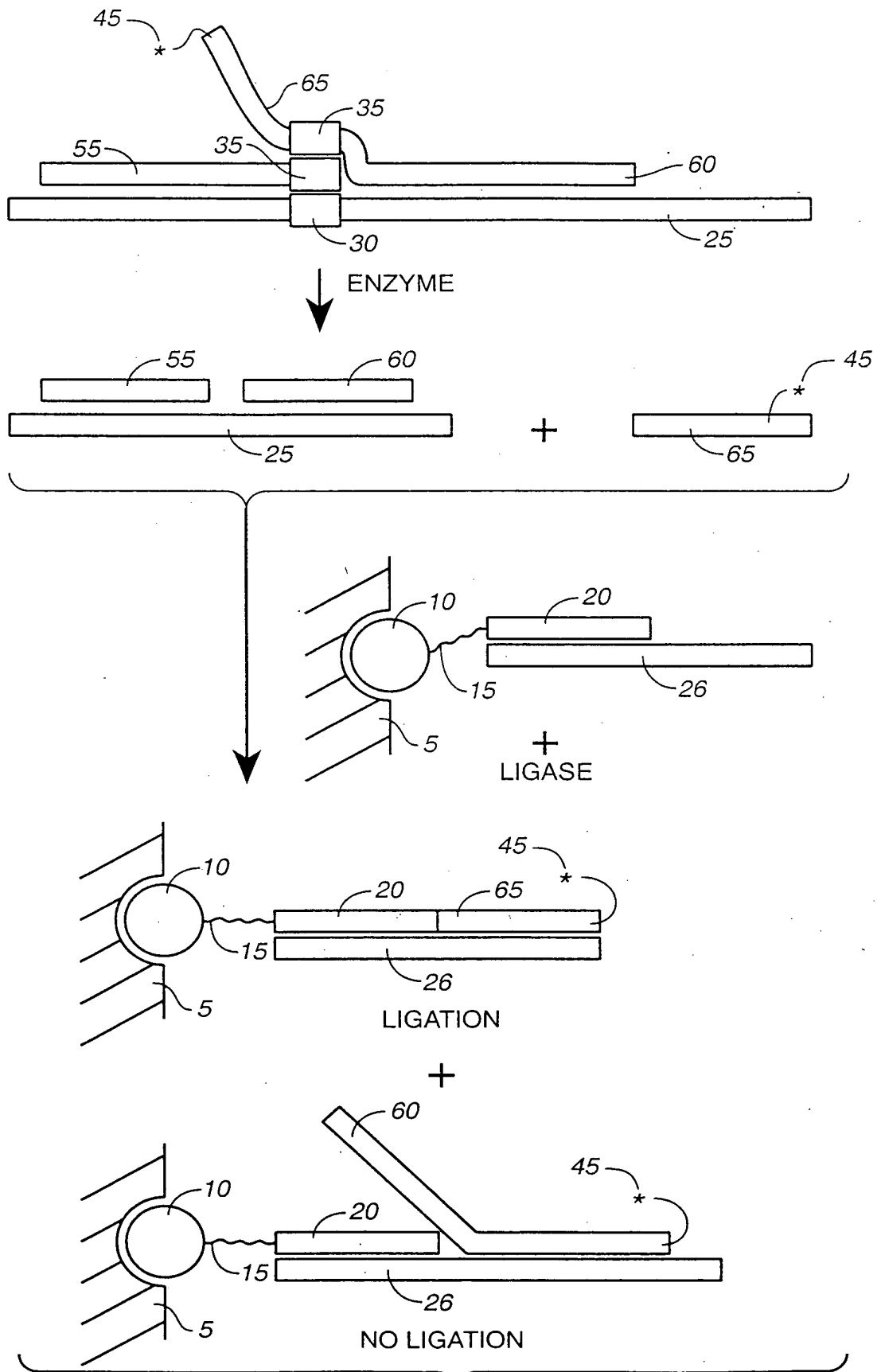


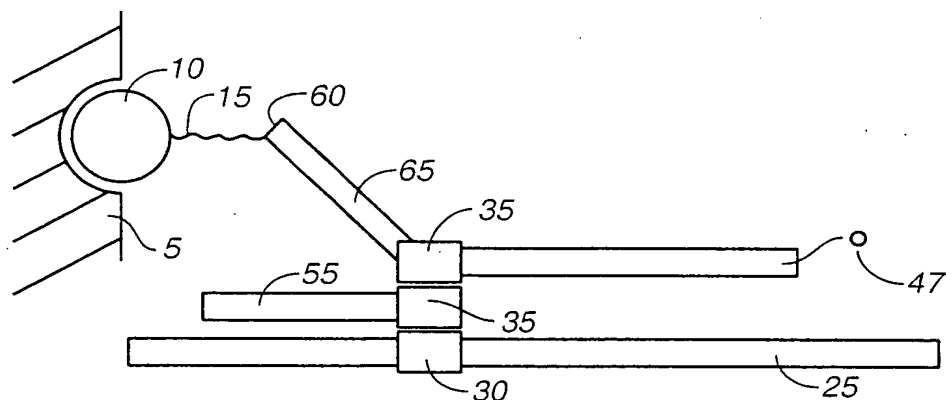
21 / 36



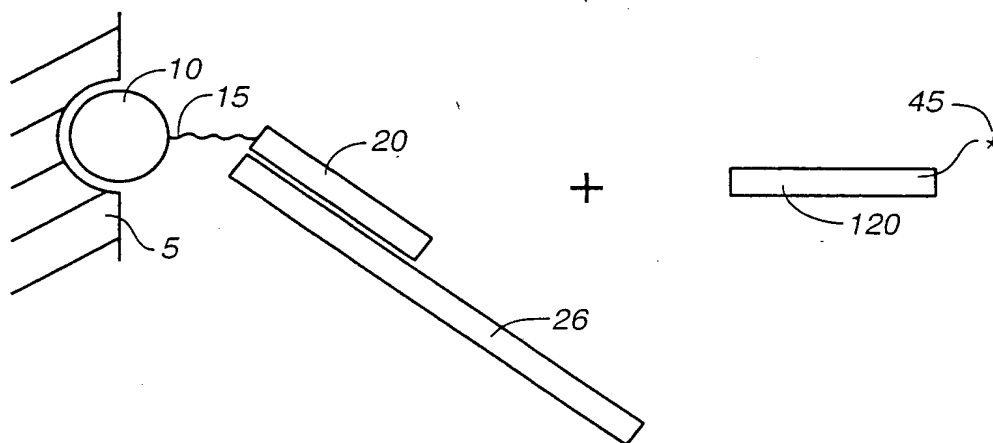
**FIG. 14C****FIG. 14D**

23 / 36

**FIG. 15A**



CLEAVAGE ENZYME
TARGET TEMPLATE



LIGATION

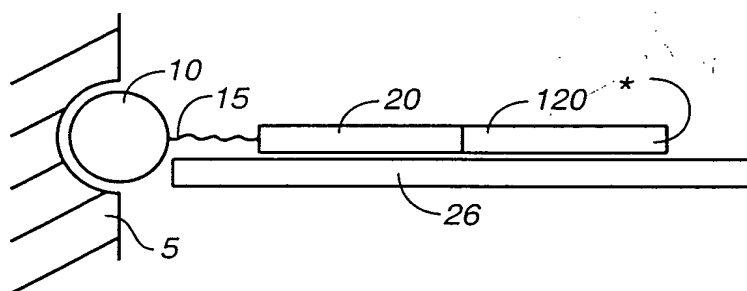
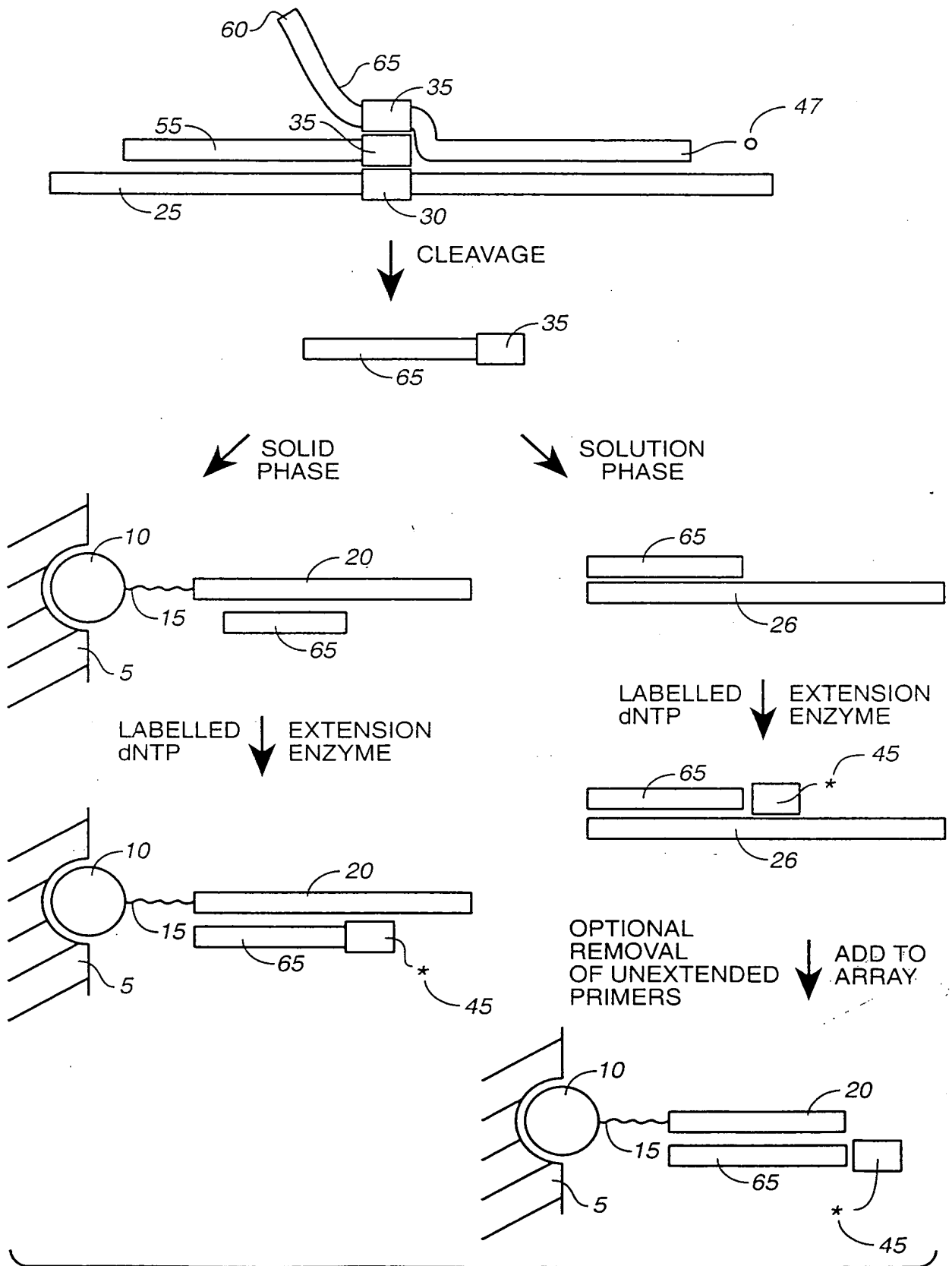
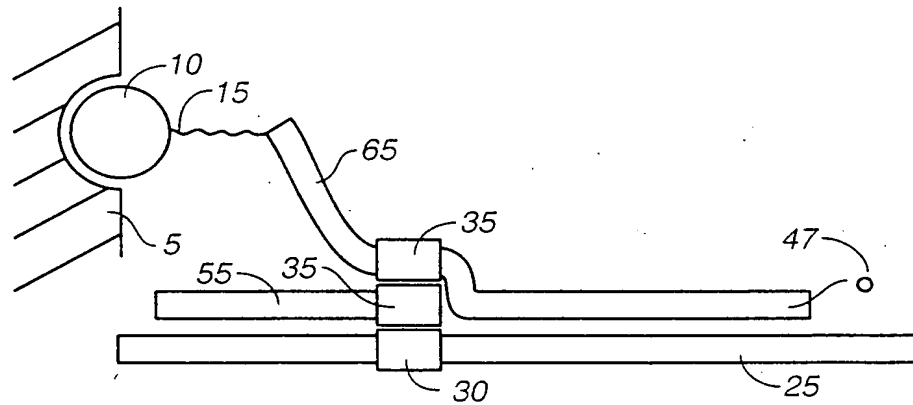


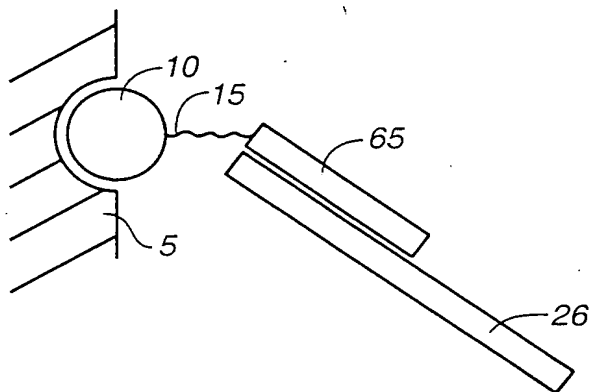
FIG. 15B

25 / 36

**FIG. 16A**



↓
CLEAVAGE
ADD TARGET TEMPLATE



↓
LABELLED
dNTP EXTENSION
 ENZYME

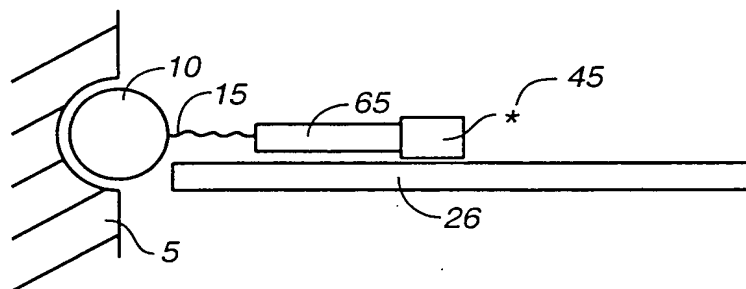
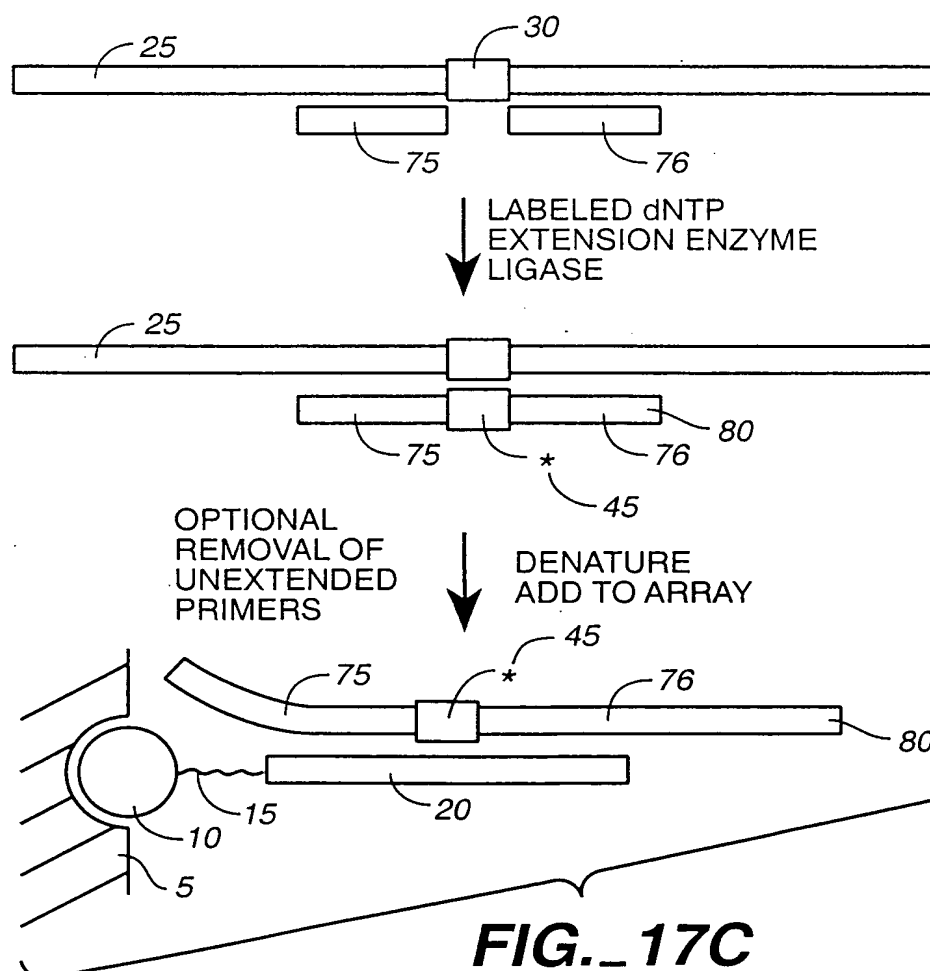
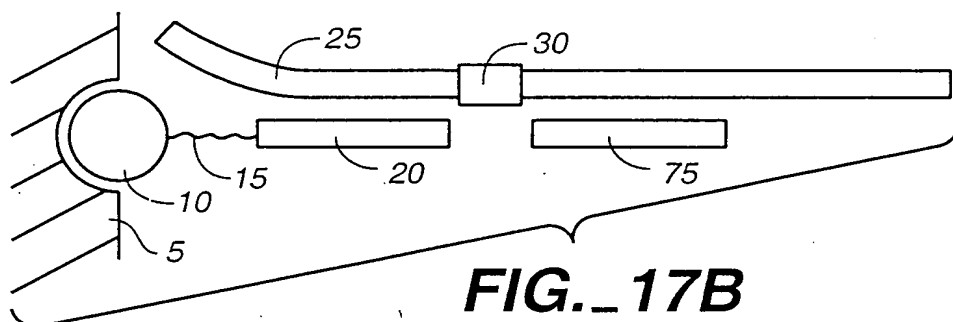
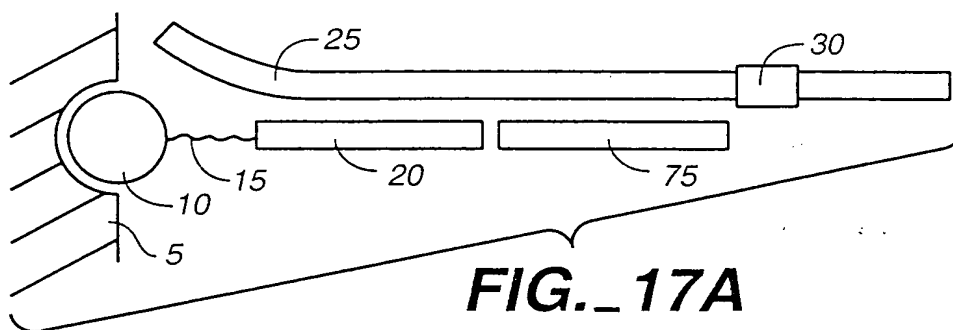
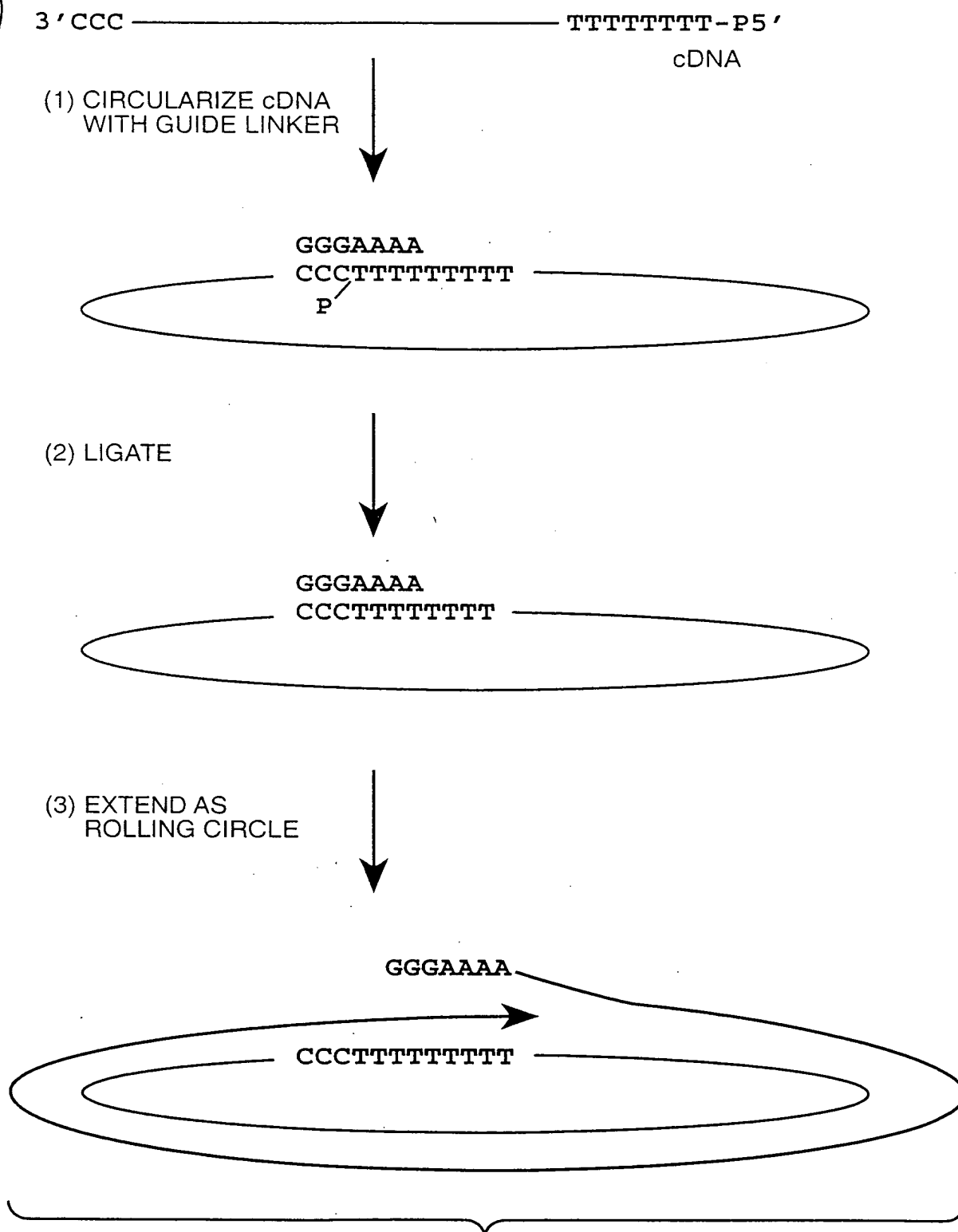


FIG. 16B



27 / 37



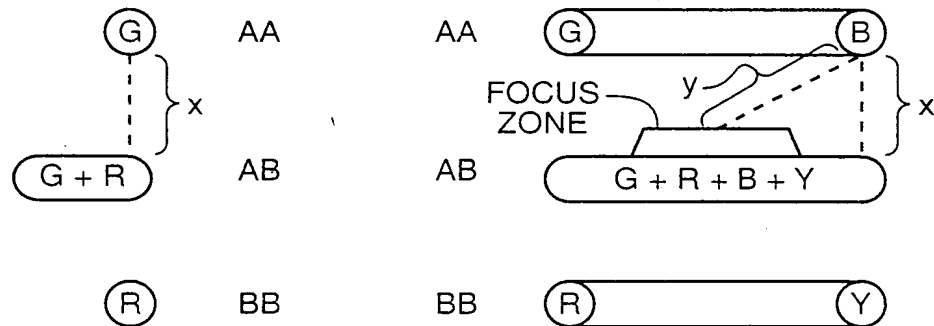
**FIG. 18**

SINGLE LABELED PROBE

GENOTYPE	SIGNAL
AA	G / G
AB	G / R
BB	R / R

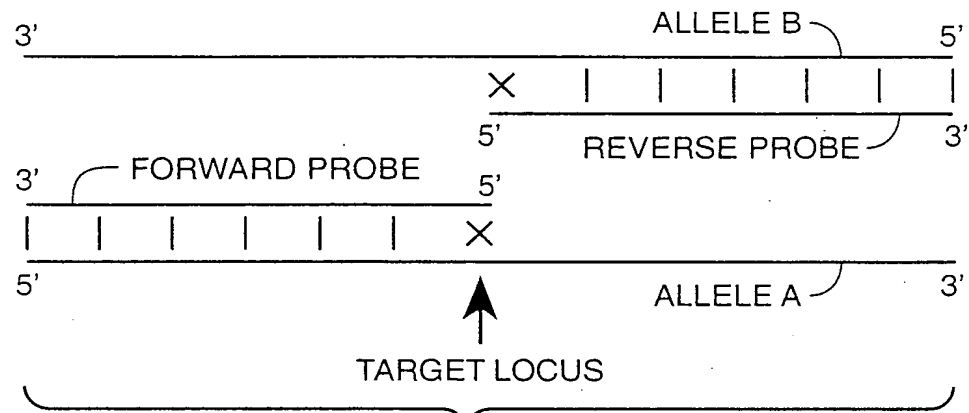
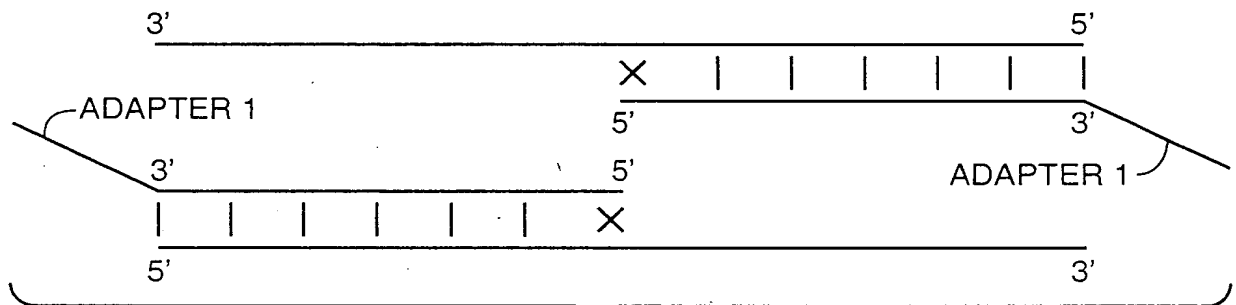
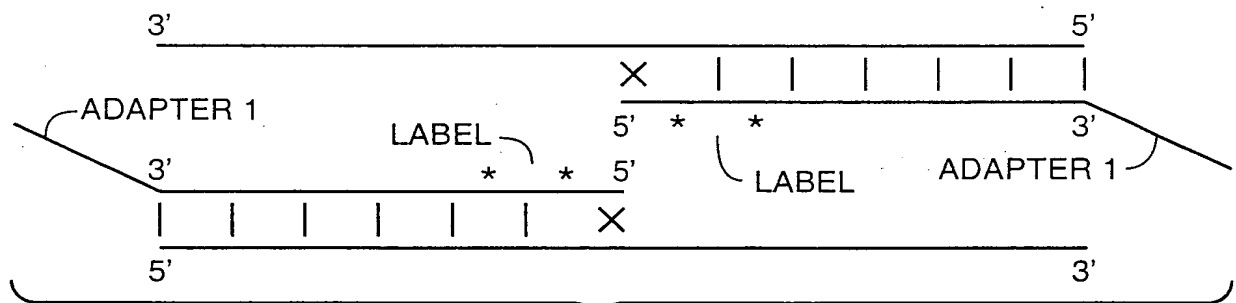
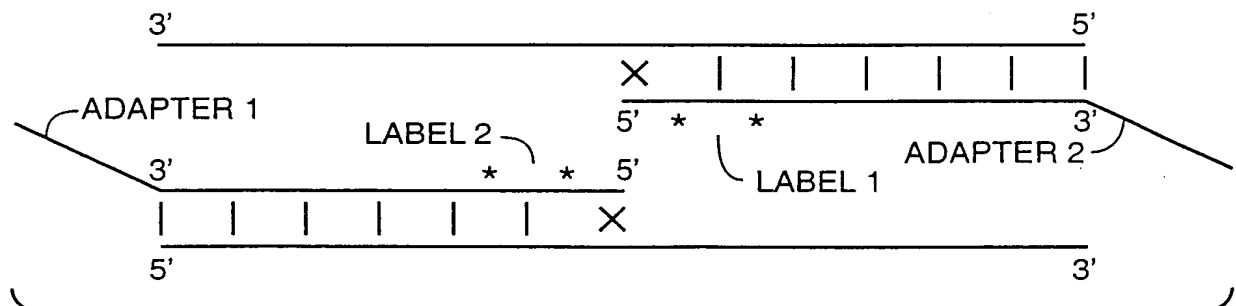
MULTI-LABELED PROBE

GENOTYPE	SIGNAL
AA	G,B / G,B
AB	G,B / R,Y
BB	R,Y / R,Y

SIGNAL RANGE

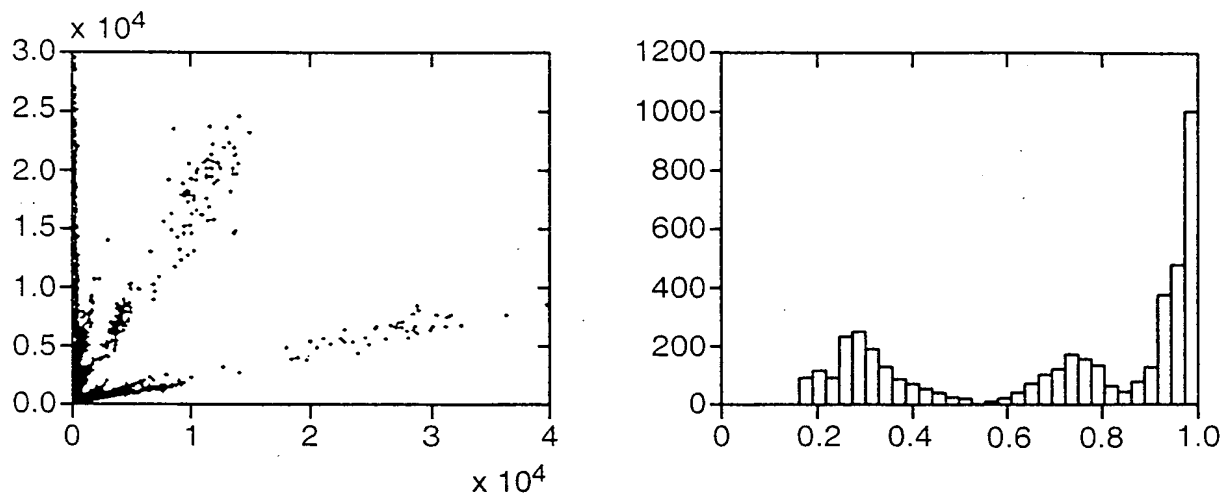
x = SINGLE LABEL DISTANCE
y = MULTI-LABEL DISTANCE

FIG. 19

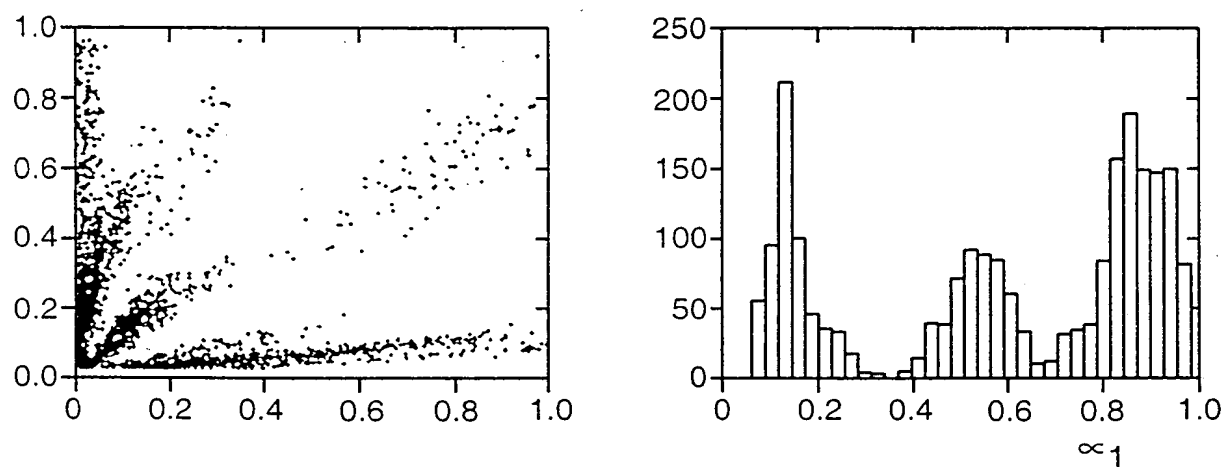
**FIG._20A****FIG._20B****FIG._20C****FIG._20D**



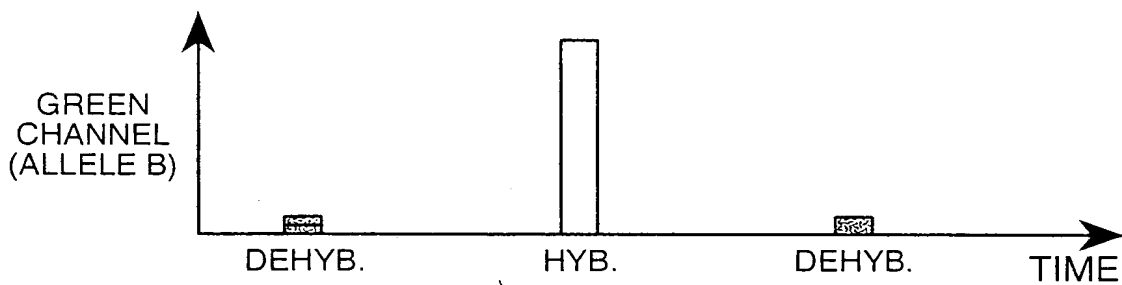
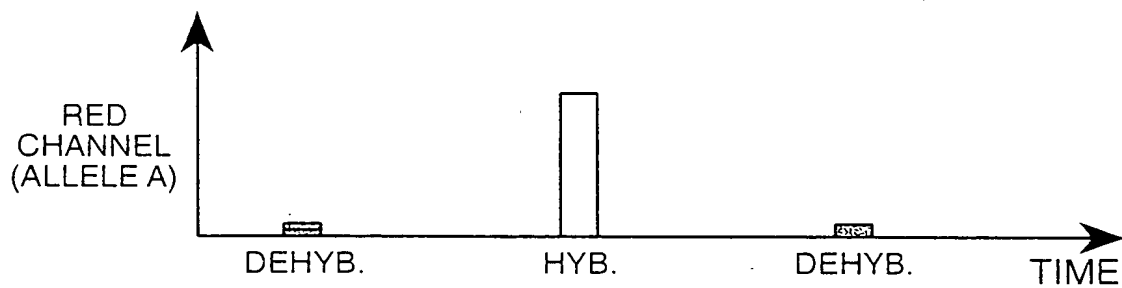
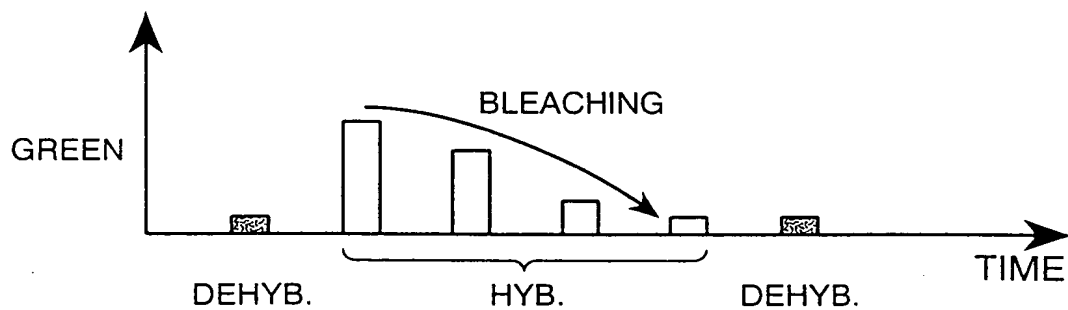
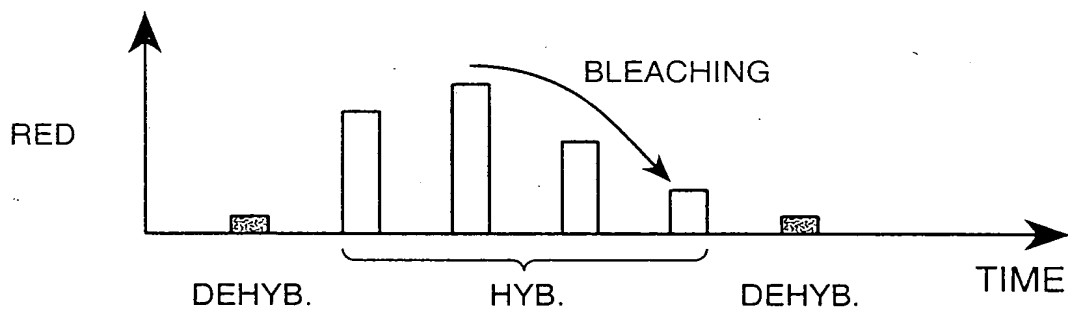
BEFORE NORMALIZATION

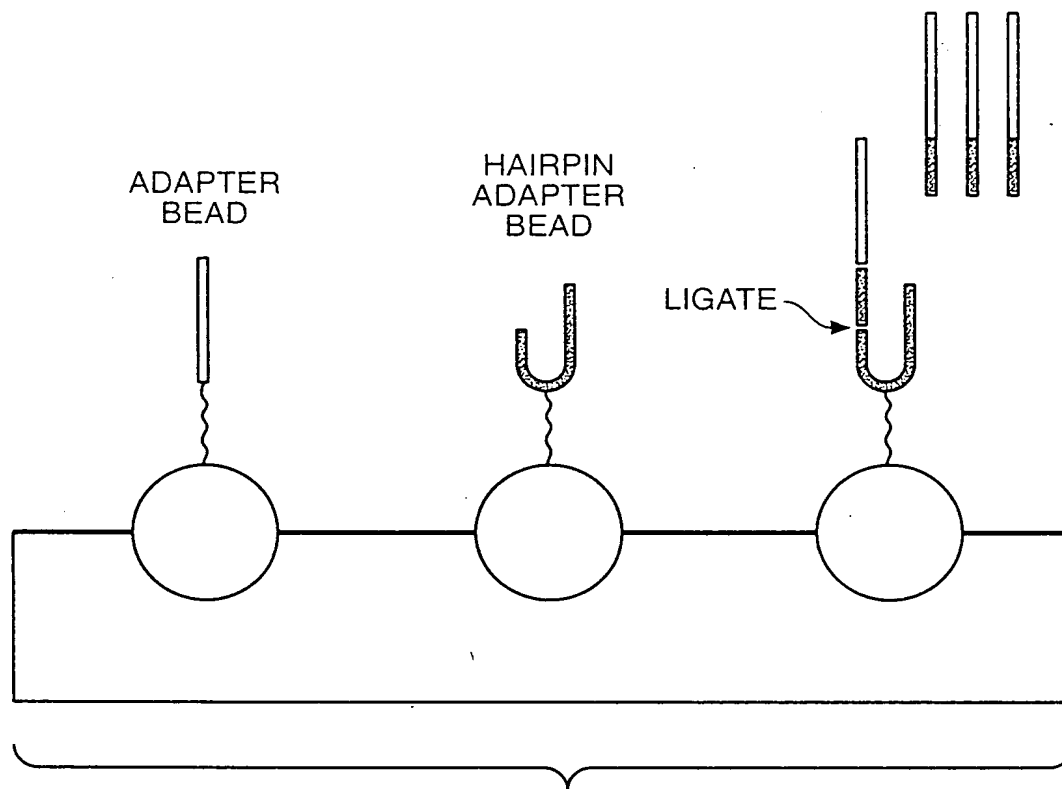


AFTER NORMALIZATION

**FIG._21**

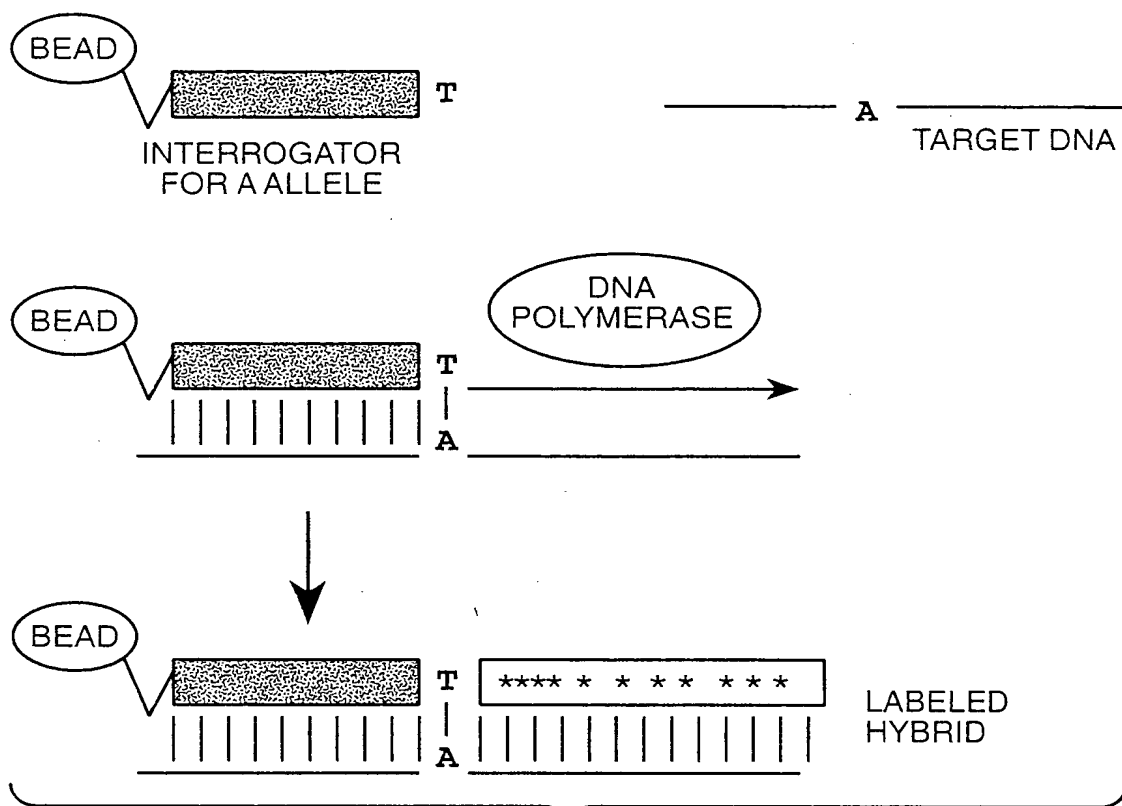
32 / 36

**FIG._22A****FIG._22B**

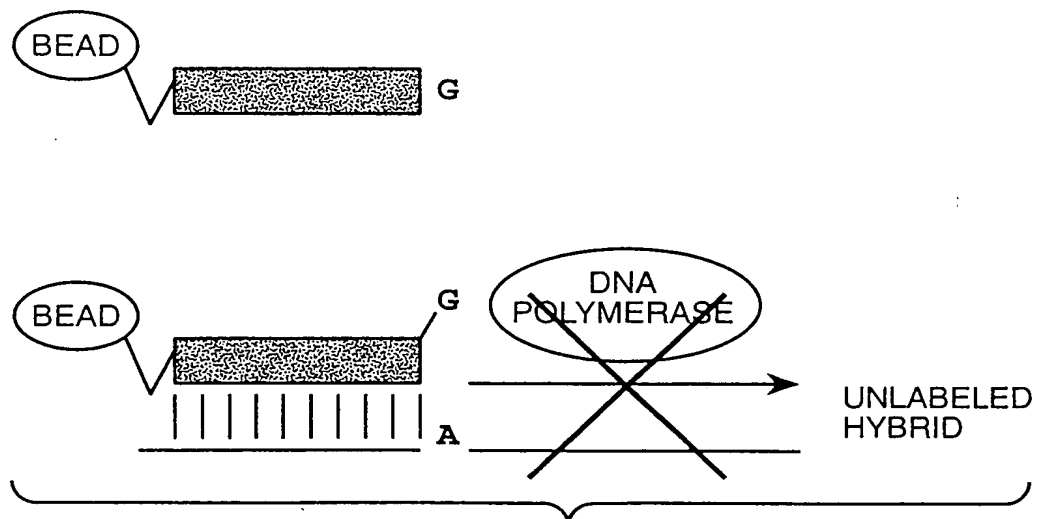
**FIG. 23**

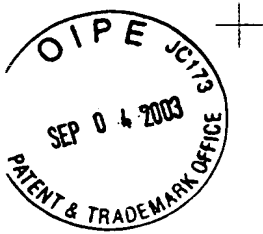


MATCH TO SNP ALLELE

**FIG._24A**

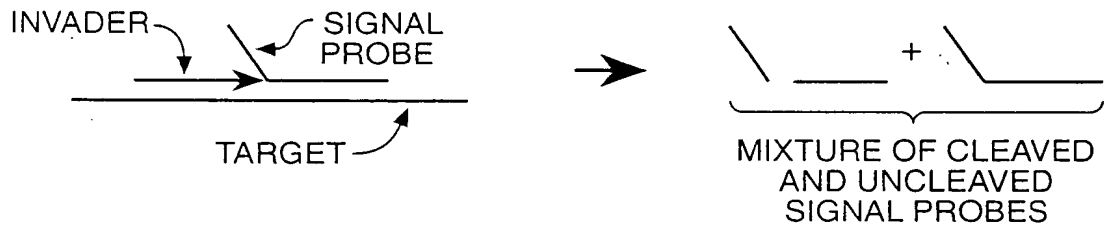
MISMATCH TO SNP ALLELE

**FIG._24B**

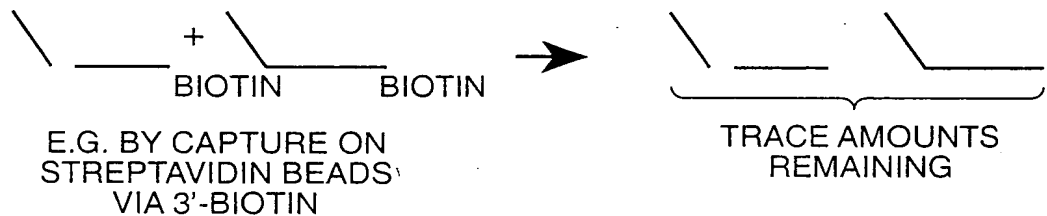


INVADER - PCR

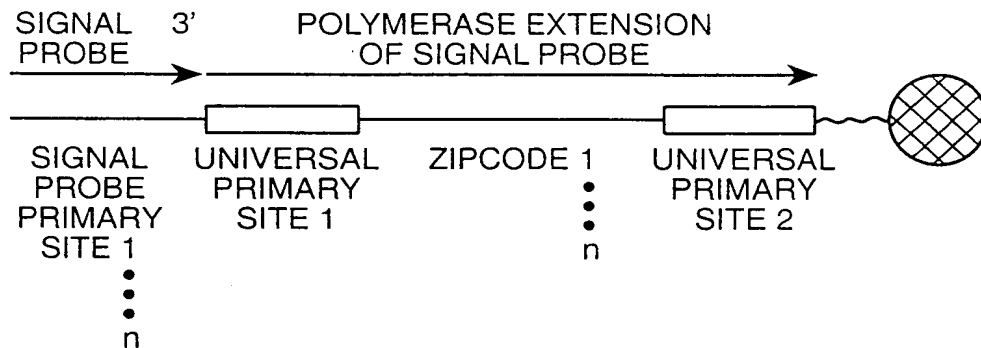
1) INVADER REACTION



2) REMOVAL OF UNCLEAVED SIGNAL PROBES



3) SIGNAL PROBE PRIMES SYNTHESIS OF AMPLICON TARGET STRAND



4) PCR AMPLIFICATION

NEWLY SYNTHESIZED TARGET STRANDS ARE DENATURED FROM TEMPLATE AND TRANSFERRED TO PCR REACTION (UNIVERSAL PRIMERS, dNT's, TAQ POLYMERASE) FOR MULTIPLEX PCR. UNIVERSAL PRIMERS ARE LABELLED E.G. WITH BIOTIN.

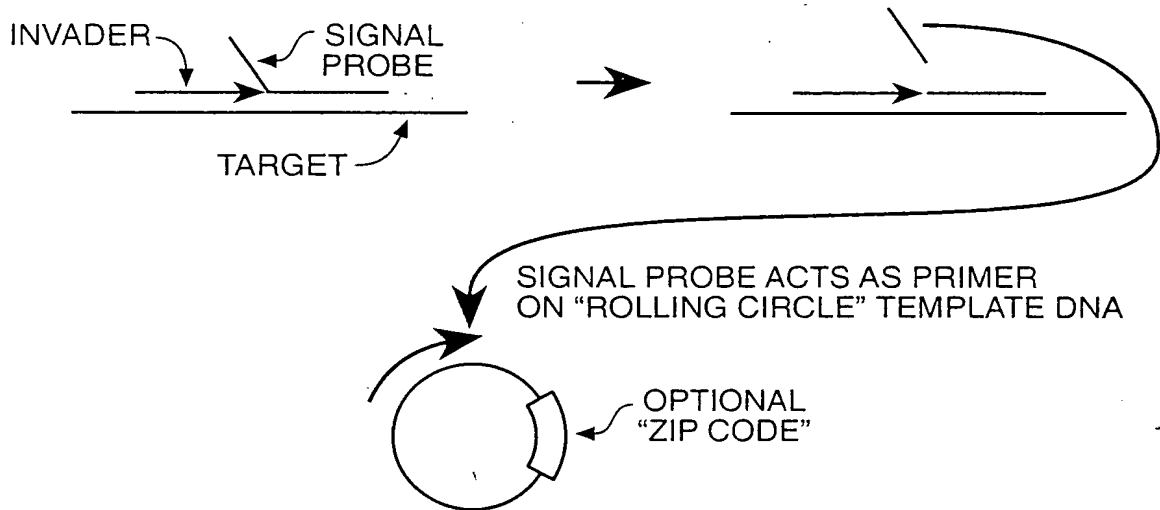
5) ARRAY HYBRIDIZATION - PCR AMPLICONS CONTAINING ZIPCODES ARE HYBRIDIZED TO ARRAY.

FIG._25

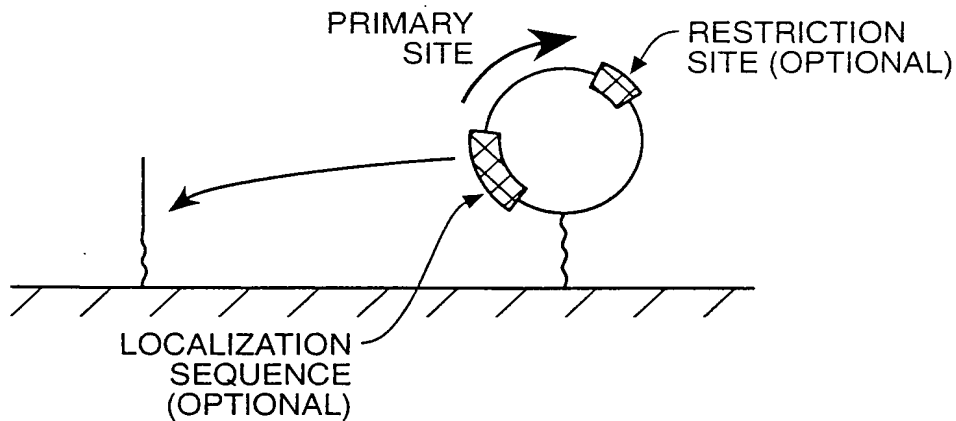


INVADER - ROLLING CIRCLE

1) INVADER REACTION



SOLID PHASE VERSION:



ROLLING CIRCLE TEMPLATE IS TETHERED TO SURFACE E.G. TO LOCALIZED "FEATURES" IN ARRAY FORMAT, OR TO BEADS.

ROLLING CIRCLE PRODUCTS CAN BE LOCALIZED E.G. BY HYBRIDIZATION TO ADJACENT PROBES OR RECOVERED IN LIQUID PHASE FOR HYBRIDIZATION TO A DETECTION ARRAY, E.G. BY ENZYMATIC CLEAVAGE.

FIG._26